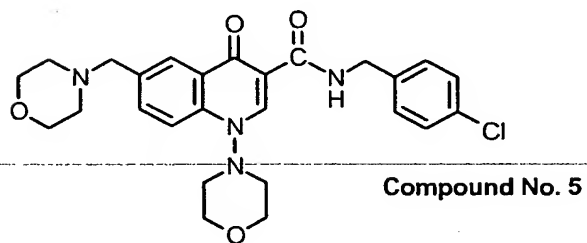
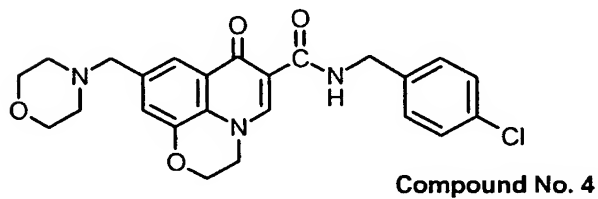
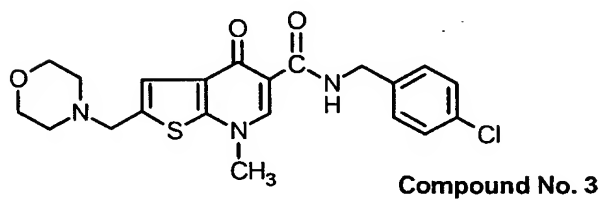
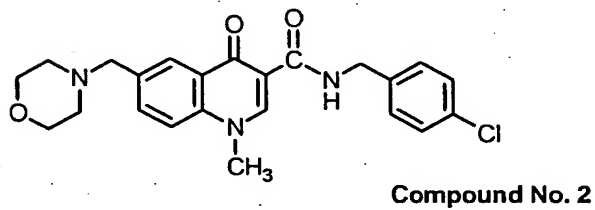
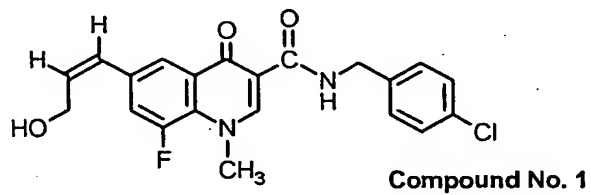
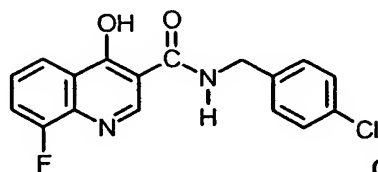


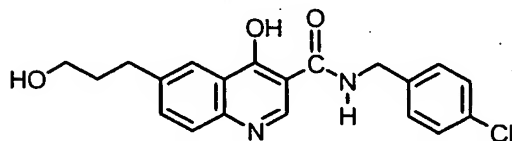
5 **Figure 1A 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds**



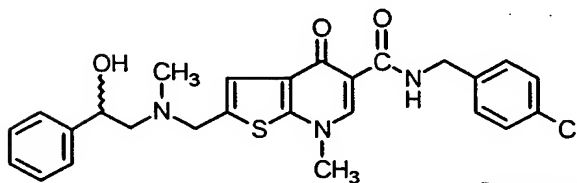
5 **Figure 1B** **4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds**



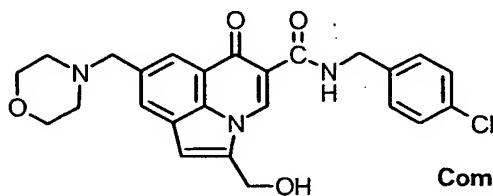
Compound No. 6



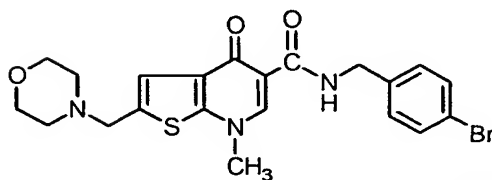
Compound No. 7



Compound No. 8

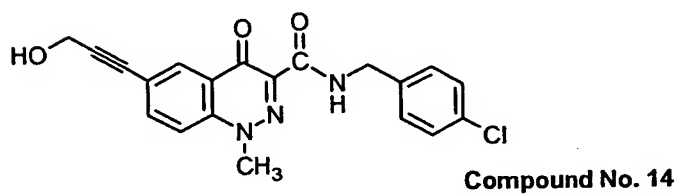
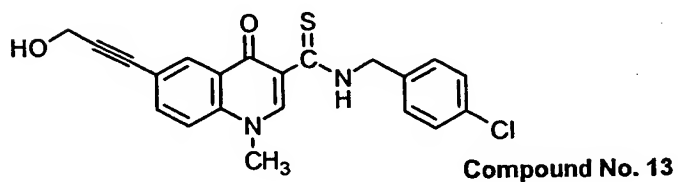
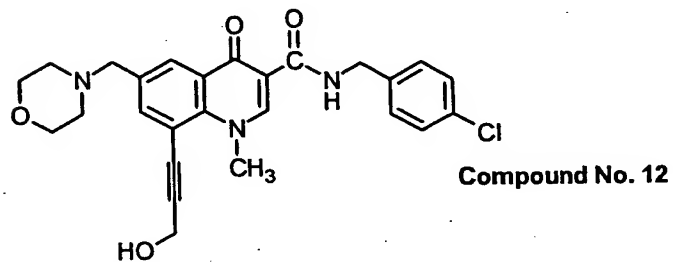
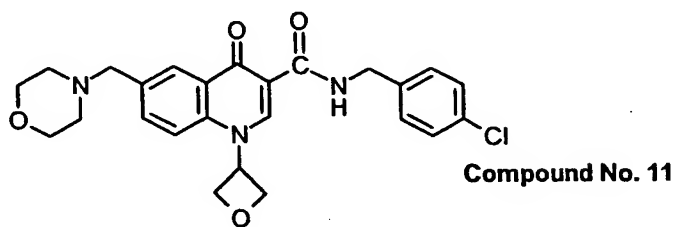


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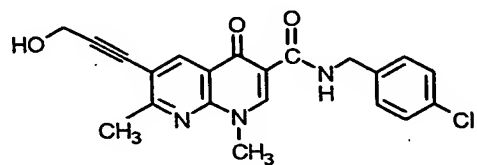


Compound No. 10

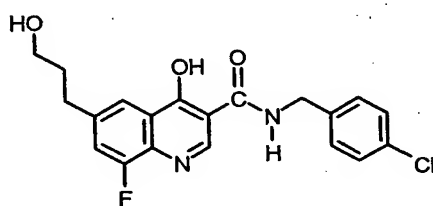
5 **Figure 1C** 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds



5 **Figure 1D** 4-HQ, 4-oxo-DHQ and 4-oxo-DHTP antiviral compounds



Compound No.15



Compound 17

Figure 4A Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Alligned by Amino Acid Homology*

| | | | | | | | |
|----|-------------|------------|------------|------------|-------------|------------|------|
| 5 | HSV2-MS | MFCAAGGPTS | PGGKSAARAA | SGFFAPHNPR | GATQTAPPPC | RRQNFYNPHL | -50 |
| | HSV2-186 | MFCAAGGPAS | PGGKSAARAA | SGFFAPHNPR | GATQTAPPPC | RRQNFYNPHL | -50 |
| | HSV1-Kos | MFSGGGGPLS | PGGKSAARAA | SGFFAPAGPR | GAGR.GPPPC | LRQNFYNPYL | -49 |
| | HSV1-Patton | MFSGGGGPLS | PGGKSAARAA | SGFFAPAGPR | GAGR.GPPPC | LRQNFYNPYL | -49 |
| | HSV1-DJL | MFSGGGGPLS | PGGKSAARAA | SGFFAPAGPR | GAGR.GPPPC | LRQNFYNPYL | -49 |
| | HSV1-F | MFSGGGGPLS | PGGKSAARAA | SGFFAPAGPR | GAGR.GPPPC | LRQNFYNPYL | -49 |
| 10 | HSV2-MS | AQTGTQPKAP | GPAQRHTYYS | ECDEFRIAP | RSLDEDAPAE | QRTGVHDGRL | -100 |
| | HSV2-186 | AQTGTQPKAP | GPAQRHTYYS | ECDEFRIAP | RSLDEDAPAE | QRTGVHDGRL | -100 |
| | HSV1-Kos | APVGTQQKPT | GPTQRHTYYS | ECDEFRIAP | RVLDEDAPPE | KRAGVHDGHL | -99 |
| | HSV1-Patton | APVGTQQKPT | GPTQRHTYYS | ECDEFRIAP | RVLDEDAPPE | KRAGVHDGHL | -99 |
| | HSV1-DJL | APVGTQQKPT | GPTQRHTYYS | ECDEFRIAP | RVLDEDAPPE | KRAGVHDGHL | -99 |
| | HSV1-F | APVGTQQKPT | GPTQRHTYYS | ECDEFRIAP | RVLDEDAPPE | KRAGVHDGHL | -99 |
| 20 | HSV2-MS | RRAPKVYCGG | DERDVLRVGP | EGFWPRRLRL | WGGADHAPKG | FDPTVTVFHV | -150 |
| | HSV2-186 | RRAPKVYCGG | DERDVLRVGP | EGFWPRRLRL | WGGADHAPEG | FDPTVTVFHV | -150 |
| | HSV-Kos | KRAPKVYCGG | DERDVLRVGS | GGFWPRRSRL | WGGVDHAPAG | FNPTVTVFHV | -149 |
| | HSV1-Patton | KRAPKVYCGG | DERDVLRVGS | GGFWPRRSRL | WGGVDHAPAG | FNPTVTVFHV | -149 |
| | HSV1-DJL | KRAPKVYCGG | DERDVLRVGS | GGFWPRRSRL | WGGVDHAPAG | FNPTVTVFHV | -149 |
| | HSV1-F | KRAPKVYCGG | DERDVLRVGS | GGFWPRRSRL | WGGVDHAPAG | FNPTVTVFHV | -149 |
| 25 | HSV2-MS | YDILEHVEHA | YSMRAAQLHE | RFMDAITPAG | TVITLLGLTP | EGHRVAVHVY | -200 |
| | HSV2-186 | YDILEHVEHA | YSMRAAQLHE | RFMDAITPAG | TVITLLGLTP | EGHRVAVHVY | -200 |
| | HSV-Kos | YDILENVEHA | YGMRAAQFHA | RFMDAITPTG | TVITLLGLTP | EGHRVAVHVY | -199 |
| | HSV1-Patton | YDILENVEHA | YGMRAAQFHA | RFMDAITPTG | TVITLLGLTP | EGHRVAVHVY | -199 |
| | HSV1-DJL | YDILENVEHA | YGMRAAQFHA | RFMDAITPTG | TVITLLGLTP | EGHRVAVHVY | -199 |
| | HSV1-F | YDILENVEHA | YGMRAAQFHA | RFMDAITPTG | TVITLLGLTP | EGHRVAVHVY | -199 |
| 30 | HSV2-MS | GTRQYFYMNK | AEVDRHLQCR | APRDLCELA | AALRESPGAS | FRGISADHFE | -250 |
| | HSV2-186 | GTRQYFYMNK | AEVDRHLQCR | APRDLCELA | AALRESPGAS | FRGISADHFE | -250 |
| | HSV-Kos | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV1-Patton | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV1-DJL | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV1-F | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| 35 | HSV2-MS | GTRQYFYMNK | AEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV2-186 | GTRQYFYMNK | AEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV-Kos | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV1-Patton | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV1-DJL | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| | HSV1-F | GTRQYFYMNK | EEVDRHLQCR | APRDLCEMA | AALRESPGAS | FRGISADHFE | -249 |
| 40 | HSV2-MS | AEVVERADVY | YYETRPTLYY | RVFVRSGRAL | AYLCDNF CPA | IRKYEGGVDA | -300 |
| | HSV2-186 | AEVVERADVY | YYETRPTLYY | RVFVRSGRAL | AYLCDNF CPA | IRKYEGGVDA | -300 |
| | HSV-Kos | AEVVERTDVY | YYETRPALFY | RVYVRSGRVL | SYLCDNF CPA | IKKYEGGVDA | -299 |
| | HSV1-Patton | AEVVERTDVY | YYETRPALFY | RVYVRSGRVL | SYLCDNF CPA | IKKYEGGVDA | -299 |
| | HSV1-DJL | AEVVERTDVY | YYETRPALFY | RVYVRSGRVL | SYLCDNF CPA | IKKYEGGVDA | -299 |
| | HSV1-F | AEVVERTDVY | YYETRPALFY | RVYVRSGRVL | SYLCDNF CPA | IKKYEGGVDA | -299 |
| 45 | HSV2-MS | TTRFILDNPG | FVTFGWYRLK | PGRGNAPAQP | RPPTAFGTSS | DVEFNCTADN | -350 |
| | HSV2-186 | TTRFILDNPG | FVTFGWYRLK | PGRGNAPAQP | RPPTAFGTSS | DVEFNCTADN | -350 |
| | HSV-Kos | TTRFILDNPG | FVTFGWYRLK | PGRNNTLAQP | RAPMAFGTSS | DVEFNCTADN | -349 |
| | HSV1-Patton | TTRFILDNPG | FVTFGWYRLK | PGRNNTLAQP | RAPMAFGTSS | DVEFNCTADN | -349 |
| | HSV1-DJL | TTRFILDNPG | FVTFGWYRLK | PGRNNTLAQP | RAPMAFGTSS | DVEFNCTADN | -349 |
| | HSV1-F | TTRFILDNPG | FVTFGWYRLK | PGRNNTLAQP | RAPMAFGTSS | DVEFNCTADN | -349 |
| 50 | HSV2-MS | LAVEGAMCDL | PAYKLMCFDI | ECKAGGEDEL | AFPVAERPED | LVIQISCLLY | -400 |
| | HSV2-186 | LAVEGAMCDL | PAYKLMCFDI | ECKAGGEDEL | AFPVAERPED | LVIQISCLLY | -400 |
| | HSV-Kos | LAIEGGMSDL | PAYKLMCFDI | ECKAGGEDEL | AFPVAGHPED | LVIQISCLLY | -399 |
| | HSV1-Patton | LAIEGGMSDL | PAYKLMCFDI | ECKAGGEDEL | AFPVAGHPED | LVIQISCLLY | -399 |
| | HSV1-DJL | LAIEGGMSDL | PAYKLMCFDI | ECKAGGEDEL | AFPVAGHPED | LVIQISCLLY | -399 |
| | HSV1-F | LAIEGGMSDL | PAYKLMCFDI | ECKAGGEDEL | AFPVAGHPED | LVIQISCLLY | -399 |
| 55 | HSV2-MS | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |
| | HSV2-186 | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |
| | HSV-Kos | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |
| | HSV1-Patton | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |
| | HSV1-DJL | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |
| | HSV1-F | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |

Figure 4B Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Aligned by Amino Acid Homology*

| | | | | | | | |
|----|-------------|-------------|-------------|-------------|------------|------------|------|
| 5 | HSV2-186 | DLSTTALEHI | LLFSLGSCDL | PESHLSDLAS | RGLPAPVVLE | FDSEFEMLLA | -450 |
| | HSV-Kos | DLSTTALEHV | LLFSLGSCDL | PESHLNELAA | RGLPTPVVLE | FDSEFEMLLA | -449 |
| | HSV1-Patton | DLSTTALEHV | LLFSLGSCDL | PESHLNELAA | RGLPTPVVLE | FDSEFEMLLA | -449 |
| | HSV1-DJL | DLSTTALEHV | LLFSLGSCDL | PESHLNELAA | RGLPTPVVLE | FDSEFEMLLA | -449 |
| | HSV1-F | DLSTTALEHV | LLFSLGSCDL | PESHLNELAA | RGLPTPVVLE | FDSEFEMLLA | -449 |
| 10 | HSV2-MS | FMTFVKQYGP | EFVTGYNIIN | FDWPFVLTKL | TEIYKVPLDG | YGRMNGRGVF | -500 |
| | HSV2-186 | FMTFVKQYGP | EFVTGYNIIN | FDWPFVLTKL | TEIYKVPLDG | YGRMNGRGVF | -500 |
| | HSV-Kos | FMTLVKQYGP | EFVTGYNIIN | FDWPFLLAKL | TDIYKVPLDG | YGRMNGRGVF | -499 |
| | HSV1-Patton | FMTLVKQYGP | EFVTGYNIIN | FDWPFLLAKL | TDIYKVPLDG | YGRMNGRGVF | -499 |
| | HSV1-DJL | FMTLVKQYGP | EFVTGYNIIN | FDWPFLLAKL | TDIYKVPLDG | YGRMNGRGVF | -499 |
| 15 | HSV1-F | FMTLVKQYGP | EFVTGYNIIN | FDWPFLLAKL | TDIYKVPLDG | YGRMNGRGVF | -499 |
| 20 | HSV2-MS | RVWDIGQSHF | QKRSKIKVNG | MVNIDMYGII | TDKVKLSSYK | LNAVAEAVLK | -550 |
| | HSV2-186 | RVWDIGQSHF | QKRSKIKVNG | MVNIDMYGII | TDKVKLSSYK | LNAVAEAVLK | -550 |
| | HSV-Kos | RVWDIGQSHF | QKRSKIKVNG | MVNIDMYGII | TDKIKLSSYK | LNAVAEAVLK | -549 |
| | HSV1-Patton | RVWDIGQSHF | QKRSKIKVNG | MVNIDMYGII | TDKIKLSSYK | LNAVAEAVLK | -549 |
| | HSV1-DJL | RVWDIGQSHF | QKRSKIKVNG | MVNIDMYGII | TDKIKLSSYK | LNAVAEAVLK | -549 |
| | HSV1-F | RVWDIGQSHF | QKRSKIKVNG | MVNIDMYGII | TDKIKLSSYK | LNAVAEAVLK | -549 |
| 25 | HSV2-MS | DKKKDLSYRD | IPAYYASGPA | QRGVIGEYCV | QDSLLVGQLF | FKFLPHLELS | -600 |
| | HSV2-186 | DKKKDLSYRD | IPAYYASGPA | QRGVIGEYCV | QDSLLVGQLF | FKFLPHLELS | -600 |
| | HSV-Kos | DKKKDLSYRD | IPAYYAAGPA | QRGVIGEYCI | QDSLLVGQLF | FKFLPHLELS | -599 |
| | HSV1-Patton | DKKKDLSYRD | IPAYYAAGPA | QRGVIGEYCI | QDSLLVGQLF | FKFLPHLELS | -599 |
| | HSV1-DJL | DKKKDLSYRD | IPAYYAAGPA | QRGVIGEYCI | QDSLLVGQLF | FKFLPHLELS | -599 |
| | HSV1-F | DKKKDLSYRD | IPAYYAAGPA | QRGVIGEYCI | QDSLLVGQLF | FKFLPHLELS | -599 |
| 30 | HSV2-MS | AVARLAGINI | TRTIYDQQI | RVFTCLLRLA | GQKGFILPDT | QGRFRGLDKE | -650 |
| | HSV2-186 | AVARLAGINI | TRTIYDQQI | RVFTCLLRLA | GQKGFILPDT | QGRFRGLDKE | -650 |
| | HSV-Kos | AVARLAGINI | TRTIYDQQI | RVFTCLLRLA | DQKGFILPDT | QGRFRGAGGE | -649 |
| | HSV1-Patton | AVARLAGINI | TRTIYDQQI | RVFTCLLRLA | DQKGFILPDT | QGRFRGAGGE | -649 |
| | HSV1-DJL | AVARLAGINI | TRTIYDQQI | RVFTCLLRLA | DQKGFILPDT | QGRFRGAGGE | -649 |
| 35 | HSV1-F | AVARLAGINI | TRTIYDQQI | RVFTCLLRLA | DQKGFILPDT | QGRFRGGGGE | -649 |
| 40 | HSV2-MS | APKRPAVPRG | EGERPGDGNG | DEKDDEDE.. | DEDGDERE.E | VARETGGRHV | -697 |
| | HSV2-186 | APKRPAVPRG | EGERPGDGNG | DEKDDEDEDEG | DEDGDERE.E | VARETGGRHV | -697 |
| | HSV-Kos | APKRPAARE | DEERP..... | EEGEDEDER | EEGGGEREPE | GARETAGRHV | -694 |
| | HSV1-Patton | APKRPAARE | DEERP..... | EEGEDEDER | EEGGGEREPE | GARETAGRHV | -694 |
| | HSV1-DJL | APKRPAARE | DEERP..... | EEGEDEDER | EEGGGEREPE | GARETAGRHV | -694 |
| | HSV1-F | APKRPAARE | DEERP..... | EEGEDEDER | EEGGGEREPE | GARETAGRHV | -694 |
| 45 | HSV2-MS | GYQGARVLD | TSGFHVDPV | VDFASLYPS | IIQAHNLCFS | TLSLRPEAVA | -747 |
| | HSV2-186 | GYQGARVLD | TSGFHVDPV | VDFASLYPS | IIQAHNLCFS | TLSLRPEAVA | -749 |
| | HSV-Kos | GYQGARVLD | TSGFHVNPV | VDFASLYPS | IIQAHNLCFS | TLSLRADAVA | -744 |
| | HSV1-Patton | GYQGARVLD | ISGFHVNPV | VDFASLYPS | IIQAHNLCFS | TLSLRADAVA | -744 |
| | HSV1-DJL | GYQGARVLD | TSGFHVNPV | VDFASLYPS | IIQAHNLCFS | TLSLRADAVA | -744 |
| 50 | HSV1-F | GYQGARVLD | TSGFHVNPV | VDFASLYPS | IIQAHNLCFS | TLSLRADAVA | -744 |
| 55 | HSV2-MS | HLEADR DYLE | IEVGGRRLFF | VKAHVRESLL | SILLRDWLAM | RKQIRSRIPQ | -797 |
| | HSV2-186 | HLEADR DYLE | IEVGGRRLFF | VKAHVRESLL | SILLRDWLAM | RKQIRSRIPQ | -799 |
| | HSV-Kos | HLEAGK DYLE | IEVGGRRLFF | VKAHVRESLL | SILLRDWLAM | RKQIRSRIPQ | -794 |
| | HSV1-Patton | HLEAGK DYLE | IEVGGRRLFF | VKAHVRESLL | SILLRDWLAM | RKQIRSRIPQ | -794 |
| | HSV1-DJL | HLEAGK DYLE | IEVGGRRLFF | VKAHVRESLL | SILLRDWLAM | RKQIRSRIPQ | -794 |
| | HSV1-F | HLEAGK DYLE | IEVGGRRLFF | VKAHVRESLL | SILLRDWLAM | RKQIRSRIPQ | -794 |
| 60 | HSV2-MS | STPEEAVLLD | KQQAIAIKVVC | NSVYGFTGVQ | HGLLPCLHVA | ATVTTIGREM | -847 |
| | HSV2-186 | SPPEEAVLLD | KQQAIAIKVVC | NSVYGFTGVQ | HGLLPCLHVA | ATVTTIGREM | -849 |
| | HSV-Kos | SSPEEAVLLD | KQQAIAIKVVC | NSVYGFTGVQ | HGLLPCLHVA | ATVTTIGREM | -844 |

Figure 4C Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Aligned by Amino Acid Homology*

| | | | | | | | |
|----|-------------|------------|------------|------------|------------|------------|-------|
| 5 | HSV1-Patton | SSPEEAVLLD | KQQAAIKVVV | NSVYGFTGVQ | HGLLPCLHVA | ATVTTIGREM | -844 |
| | HSV1-DJL | SSPEEAVLLD | KQQAAIKVVV | NSVYGFTGVQ | HGLLPCLHVA | ATVTTIGREM | -844 |
| | HSV1-F | SSPEEAVLLD | KQQAAIKVVV | NSVYGFTGVQ | HGLLPCLHVA | ATVTTIGREM | -844 |
| | | | | | | | |
| 10 | HSV2-MS | LLATRAYVHA | RWAEFDQLLA | DFPEAAGMRA | PGPYSMRIIY | GDTSIFVLC | -897 |
| | HSV2-186 | LLATRAYVHA | RWAEFDQLLA | DFPEAAGMRA | PGPYSMRIIY | GDTSIFVLC | -899 |
| | HSV-Kos | LLATREYVHA | RWAAFEQLLA | DFPEADMRA | PGPYSMRIIY | GDTSIFVLC | -894 |
| | HSV1-Patton | LLATREYVHA | RWAAFEQLLA | DFPEADMRA | PGPYSMRIIY | GDTSIFVLC | -894 |
| | HSV1-DJL | LLATREYVHA | RWAAFEQLLA | DFPEADMRA | PGPYSMRIIY | GDTSIFVLC | -894 |
| | HSV1-F | LLATREYVHA | RWAAFEQLLA | DFPEADMRA | PGPYSMRIIY | GDTSIFVLC | -894 |
| | | | | | | | |
| 15 | HSV2-MS | RGLTAAGLVA | MGDKMASHIS | RALFLPPIKL | ECEKTFTKLL | LIAKKKYIGV | -947 |
| | HSV2-186 | RGLTAAGLVA | MGDKMASHIS | RALFLPPIKL | ECEKTFTKLL | LIAKKKYIGV | -949 |
| | HSV-Kos | RGLTAAGLTA | MGDKMASHIS | RALFLPPIKL | ECEKTFTKLL | LIAKKKYIGV | -944 |
| | HSV1-Patton | RGLTAAGLTA | MGDKMASHIS | RALFLPPIKL | ECEKTFTKLL | LIAKKKYIGV | -944 |
| | HSV1-DJL | RGLTAAGLTA | VGDKMASHIS | RALFLPPIKL | ECEKTFTKLL | LIAKKKYIGV | -944 |
| | HSV1-F | RGLTAAGLTA | VGDKMASHIS | RALFLSPIKL | ECEKTFTKLL | LIAKKKYIGV | -944 |
| | | | | | | | |
| 25 | HSV2-MS | ICGGKMLIKG | VDLVRKNNCA | FINRTSRALV | DLLFYDDTVS | GAAAALAERP | -997 |
| | HSV2-186 | ICGGKMLIKG | VDLVRKNNCA | FINRTSRALV | DLLFYDDTVS | GAAAALAERP | -999 |
| | HSV-Kos | IYGGKMLIKG | VDLVRKNNCA | FINRTSRALV | DLLFYDDTVS | GAAAALAERP | -994 |
| | HSV1-Patton | IYGGKMLIKG | VDLVRKNNCA | FINRTSRALV | DLLFYDDTVS | GAAAALAERP | -994 |
| | HSV1-DJL | IYGGKMLIKG | VDLVRKNNCA | FINRTSRALV | DLLFYDDTVS | GAAAALAERP | -994 |
| | HSV1-F | IYGGKMLIKG | VDLVRKNNCA | FINRTSRALV | DLLFYDDTVS | GAAAALAERP | -994 |
| | | | | | | | |
| 30 | HSV2-MS | AEEWLARPLP | EGLQAFGAVL | VDAHRRITDP | ERDIQDFVLT | AELSRHPRAY | -1047 |
| | HSV2-186 | AEEWLARPLP | EGLQAFGAVL | VDAHRRITDP | ERDIQDFVLT | AELSRHPRAY | -1049 |
| | HSV-Kos | AEEWLARPLP | EGLQAFGAVL | VDAHRRITDP | ERDIQDFVLT | AELSRHPRAY | -1044 |
| | HSV1-Patton | AEEWLARPLP | EGLQAFGAVL | VDAHRRITDP | ERDIQDFVLT | AELSRHPRAY | -1044 |
| | HSV1-DJL | AEEWLARPLP | EGLQAFGAVL | VDAHRRITDP | ERDIQDFVLT | AELSRHPRAY | -1044 |
| | HSV1-F | AEEWLARPLP | EGLQAFGAVL | VDAHRRITDP | ERDIQDFVLT | AELSRHPRAY | -1044 |
| | | | | | | | |
| 35 | HSV2-MS | TNKRLAHLTV | YYKLMARRAQ | VPSIKDRIPY | VIVAQTREVE | ETVARLAALR | -1097 |
| | HSV2-186 | TNKRLAHLTV | YYKLMARRAQ | VPSIKDRIPY | VIVAQTREVE | ETVARLAALR | -1099 |
| | HSV-Kos | TNKRLAHLTV | YYKLMARRAQ | VPSIKDRIPY | VIVAQTREVE | ETVARLAALR | -1094 |
| | HSV1-Patton | TNKRLAHLTV | YYKLMARRAQ | VPSIKDRIPY | VIVAQTREVE | ETVARLAALR | -1094 |
| | HSV1-DJL | TNKRLAHLTV | YYKLMARRAQ | VPSIKDRIPY | VIVAQTREVE | ETVARLAALR | -1094 |
| | HSV1-F | TNKRLAHLTV | YYKLMARRAQ | VPSIKDRIPY | VIVAQTREVE | ETVARLAALR | -1094 |
| | | | | | | | |
| 45 | HSV2-MS | ELDAAAPGDE | PAPPAALPSP | AKRPRETPSH | ADPPGGASKP | RKLLVSELAE | -1147 |
| | HSV2-186 | ELDAAAPGDE | PAPPAALPSP | AKRPRETPSH | ADPPGGASKP | RKLLVSELAE | -1149 |
| | HSV-Kos | ELDAAAPGDE | PAPPAALPSP | AKRPRETPSH | ADPPGGASKP | RKLLVSELAE | -1144 |
| | HSV1-Patton | ELDAAAPGDE | PAPPAALPSP | AKRPRETPSP | ADPPGGASKP | RKLLVSELAE | -1144 |
| | HSV1-DJL | ELDAAAPGDE | PAPPAALPSP | AKRPRETPSP | ADPPGGASKP | RKLLVSELAE | -1144 |
| | HSV1-F | ELDAAAPGDE | PAPPAALPSP | AKRPRETPLH | ADPPGGASKP | RKLLVSELAE | -1144 |
| | | | | | | | |
| 50 | HSV2-MS | DPGYAIARGV | PLNTDYYFSH | LLGAACVTFK | ALFGNNAKIT | ESLLKRFIPE | -1197 |
| | HSV2-186 | DPGYAIARGV | PLNTDYYFSH | LLGAACVTFK | ALFGNNAKIT | ESLLKRFIPE | -1199 |
| | HSV-Kos | DPAYAIAGHV | ALNTDYYFSH | LLGAACVTFK | ALFGNNAKIT | ESLLKRFIPE | -1194 |
| | HSV1-Patton | DPAYAIAGHV | ALNTDYYFSH | LLGAACVTFK | ALFGNNAKIT | ESLLKRFIPE | -1194 |
| | HSV1-DJL | DPAYAIAGHV | ALNTDYYFSH | LLGAACVTFK | ALFGNNAKIT | ESLLKRFIPE | -1194 |
| | HSV1-F | DPAYAIAGHV | ALNTDYYFSH | LLGAACVTFK | ALFGNNAKIT | ESLLKRFIPE | -1194 |
| | | | | | | | |
| 60 | HSV2-MS | TWHPPDDVAA | RLRAAGFGPA | GAGATAEETR | RMLHRAFDTL | A* | -1238 |
| | HSV2-186 | TWHPPDDVAA | RLRAAGFGPA | GAGATAEETR | RMLHRAFDTL | A* | -1240 |
| | HSV-Kos | VWHPPDDVAA | RLRAAGFGAV | GAGATAEETR | RMLHRAFDTL | A* | -1235 |
| | HSV1-Patton | VWHPPDDVTA | RLRAAGFGAV | GAGATAEETR | RMLHRAFDTL | A* | -1235 |
| | HSV1-DJL | VWHPPDDVAA | RLRTAGFGAV | GAGATAEETR | RMLHRAFDTL | A* | -1235 |
| | | | | | | | |

Figure 4D Comparison of Wild type HSV-1 and HSV-2 DNA Polymerases Amino Acid Sequences Alligned by Amino Acid Homology*

5 HSV1-F VWHPPDDVAA RLRAAGFGAV GAGATAEETR RMLHRAFDTL A* -1235

*Amino acid alignment demonstrates difference in amino acid's sequences.

*The gaps "....." indicate missing amino acids relative to other stanins.

*Wild HSV2-MS is listed as SEQ. ID NO 14.

10 *Wild HSV2-186 is listed as SEQ. ID NO 15.

*Wild HSV-Kos is listed as SEQ. ID NO 16.

*Wild HSV1-Patton is listed as SEQ. ID NO 17.

*Wild HSV1-DJL is listed as SEQ. ID NO 18.

*Wild HSV1-F is listed as SEQ. ID NO 19.

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5 **Figure 5A DNA and amino acid sequence list****SEQ. ID. NO. 1** DNA sequence of DNA polymerase gene for HSV2-MS-M1

1 ATGTTTGTG CCGCGGGCGG CCCGACTTCC CCCGGGGGGA AGTCGGCGGC
 10 51 TCGGGCGGCG TCTGGGTTTT TTGCCCCCA CAACCCCGG GGAGCCACCC
 101 AGACGGCACC GCCGCCTTGC CGCCGGCAGA ACTTCTACAA CCCCCACCTC
 15 151 GCTCAGACCG GAACGCAGCC AAAGGCCCCC GGGCGGCTC AGCGCCATAC
 201 GTACTACAGC GAGTGCGACG AATTTCGATT TATCGCCCCG CGTTCGCTGG
 251 ACGAGGACGC CCCCGCGGAG CAGCGCACCG GGGTCCACGA CGGCCGCCTC
 20 301 CGGCGCGCCC CTAAGGTGTA CTGCGGGGGG GACGAGCGCG ACGTCCTCCG
 351 CGTGGGCCCG GAGGGCTTCT GGCCGCGTCG CTTGCGCCTG TGGGGCGGTG
 25 401 CGGACCATGC CCCCAAGGGG TTCGACCCCA CCGTCACCGT CTTCCACGTG
 451 TACGACATCC TGGAGCACGT GGAACACGCG TACAGCATGC GCGCCGCCCA
 501 GCTCCACGAG CGATTTATGG ACGCCATCAC GCCCGCCGGG ACGTCATCA
 30 551 CGCTTCTGGG TCTGACCCCC GAAGGCCATC GCGTCGCCGT TCACGTCTAC
 601 GGCACGCGGC AGTACTTTTA CATGAACAAG GCGGAGGTGG ATCGGCACCT
 35 651 GCAGTGCCGT GCCCCGCGCG ATCTCTGCGA GCGCCTGGCG GCGGCCCTGC
 701 GCGAGTCGCC GGGGGCGTCG TTCCGCGGCA TCTCCGCGGA CCACTTOGAG
 751 GCGGAGGTGG TGGAGCGCGC CGACGTGTAC TATTACGAAA CGCGCCCGAC
 40 801 CCTGTACTAC CGCGTCTTCG TCGAAGCGG GCGCGCGCTG GCCTACCTGT
 851 GCGACAACCT TTGCCCCGCG ATCAGGAAGT ACGAGGGGGG CGTCGACGCC
 45 901 ACCACCGGT TTATCCTGGA CAACCCGGGG TTTGTACCT TCGGCTGGTA
 951 CCGCCTCAAG CCCGGCCGCG GGAACGCGCC GGCCCAACCG CGCCCCCGA
 1001 CGGCGTTCGG AACCTCGAGC GACGTCGAGT TTAACGAC GCGGACAAC
 50 1051 CTGGCCGTCG AGGGGGCCAT GTGTGACCTG CCGGCCTACA AGCTCATGTG
 1101 CTTGATATC GAATGCAAGG CCGGGGGGGA GGACGAGCTG GCCTTTCGG
 55 1151 TCGCGGAACG CCCGGAAGAC CTCGTCATCC AGATCTCCTG TCTGCTCTAC
 1201 GACCTGTCCA CCACCGCCCT CGAGCACATC CTCCTGTTTT CGCTCGGATC

5 **Figure 5B DNA and amino acid sequence list**

1251 CTGCGACCTC CCCGAGTCCC ACCTCAGCGA TCTCGCCTCC AGGGGCCTGC
 1301 CGGCCCCCGT CGTCCTGGAG TTTGACAGCG AATTCGAGAT GCTGCTGGCC
 10 1351 TTCATGACCT TCGTCAAGCA GTACGGCCCC GAGTTCGTGA CCGGGTACAA
 1401 CATCATCAAC TTCGACTGGC CCTTCGTCTT GACCAAGCTG ACGGAGATCT
 15 1451 ACAAGGTCCC GCTCGACGGG TACGGGCGCA TGAACGGCCG GGGTGTGTTC
 1501 CGCGTGTGGG ACATCGGCCA GAGCCACTTT CAGAAGCGCA GCAAGATCAA
 1551 GGTGAACGGG ATGGTGAACA TCGACATGTA CGGCATCATC ACGGACAAGG
 20 1601 TCAAACCTCT CAGCTACAAG CTGAACGCCG TCGCCGAGGC CGTCTTGAAG
 1651 GACAAGAAGA AGGATCTGAG CTACCGCGAC ATCCCCGCCT ACTACGCCTC
 25 1701 CGGGCCCCGG CAGCGCGGGG TGATCGGCGA GTATTGTGTG CAGGACTCGC
 1751 TGCTGGTCGG GCAGCTGTTC TTCAAGTTTC TGCCGCACCT GGAGCTTTCC
 1801 GCCGTGCGGC GCCTGGCGGG CATCAACATC ACCCGCACCA TCTACGACGG
 30 1851 CCAGCAGATC CGCGTCTTCA CGTGCTCTCT GCGCCTTGCG GGCCAGAAGG
 1901 GCTTCATCCT GCCGGACACC CAGGGGCGGT TTCGGGGCCT CGACAAGGAG
 35 1951 GCGCCCAAGC GCCCGGCCGT GCCTCGGGGG GAAGGGGAGC GGCCGGGGGA
 2001 CGGGAACGGG GACGAGGATA AGGACGACGA CGAGGACGAG GACGGGGACG
 2051 AGCGCGAGGA GGTCGCGCGC GAGACCGGGG GCGGCACGT TGGGTACCAG
 40 2101 GGGGCCCCGG TCCTCGACCC CACCTCCGGG TTTCACGTCG ACCCCGTGGT
 2151 GGTGTTTGAC TTTGCCAGCC TGTACCCAG CATCATCCAG GCCACAACC
 45 2201 TGTGCTTCAG TACGCTCTCC CTGCGGCCCC AGGCCGTCGC GCACCTGGAG
 2251 GCGGACCGGG ACTACCTGGA GATCGAGGTG GGGGGCCGAC GGCTGTTCTT
 2301 CGTGAAGGCC CACGTACGCG AGAGCCTGCT GAGCATCCTG CTGCGCGACT
 50 2351 GGCTGGCCAT GCGAAAGCAG ATCCGCTCGC GGATCCCCCA GAGCACCCCC
 2401 GAGGAGGCCG TCCTCCTCGA CAAGCAACAG GCCGCCATCA AGGTGGTGTG
 55 2451 CAACTCGGTG TACGGGTTCA CCGGGGCGCA GCACGGTCTT CTGCCCTGCC
 2501 TGCACGTGGC CGCCACCGTG ACGACCATCG GCCGCGAGAT GTCCTCGCG
 2551 ACGCGCGCGT ACGTGACGCG GCGCTGGGCG GAGTTCGATC AGCTGCTGGC
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5 **Figure 5C DNA and amino acid sequence list**

2601 CGACTTTCCG GAGGCGGCCG GCATGCGCGC CCCCGGTCCG TACTCCATGC
 2651 GCATCATCTA CGGGGACACG GACTCCATTT TCGTTTGTG CCGCGGCCTC
 10 2701 ACGGCCGCGG GCCTGGTGGC CATGGGCGAC AAGATGGCGA GCCACATCTC
 2751 GCGCGCGCTG TTCCTCCCC CGATCAAGCT CGAGTGCAGAA AAAACGTTCA
 15 2801 CCAAGCTGCT GCTCATCGCC AAGAAAAAGT ACATCGGCGT CATCTGCGGG
 2851 GGCAAGATGC TCATCAAGGG CGTGGATCTG GTGCGCAAAA ACAACTGCGC
 2901 GTTTATCAAC CGCACCTCCA GGGCCCTGGT CGACCTGCTG TTTTACGAAG
 20 2951 ATACCGTATC CGGAGCGGCC GCCGCGTTAG CCGAGCGCCC CGCAGAGGAG
 3001 TGGCTGGCGC GACCCCTGCC CGAGGGACTG CAGGCGTTCG GGGCCGTCCT
 25 3051 CGTAGACGCC CATCGGCGCA TCACCGACCC GGAGAGGGAC ATCCAGGACT
 3101 TTGTCCTCAC CGCCGAAGTG AGCAGACACC CGCGCGCGTA CACCAACAAG
 3151 CGCCTGGCCC ACCTGACGGT GTATTACAAG CTCATGGCCC GCCGCGCGCA
 30 3201 GGTCCCGTCC ATCAAGGACC GGATCCCGTA CGTGATCGTG GCCCAGACCC
 3251 GCGAGGTAGA GGAGACGGTC GCGCGGCTGG CCGCCCTCCG CGAGCTAGAC
 35 3301 GCCGCGCCC CAGGGGACGA GCCCGCCCC CCAGCGGCCC TGCCCTCCCC
 3351 GGCCAAGCGC CCCCGGGAGA CGCCGTCGCA TGCCGACCCC CCGGGAGGCG
 3401 CGTCCAAGCC CCGCAAGCTG CTGGTGTCCG AGCTGGCGGA GGATCCCGGG
 40 3451 TACGCCATCG CCCGGGGCGT TCCGCTCAAC ACGGACTATT ACTTCTCGCA
 3501 CCTGCTGGGG GCGGCCTGCG TGACGTTCAA GGCCCTGTTT GGAAATAACG
 45 3551 CCAAGATCAC CGAGAGTCTG TTAAAGAGGT TTATTCCCGA GACGTGGCAC
 3601 CCCCCGACG ACGTGGCCGC GCGGCTCAGG GCCGCGGGGT TCGGGCCGGC
 3651 GGGGGCCGGC GCTACGGCGG AGGAAACTCG TCGAATGTTG CATAGAGCCT
 50 3701 TTGATACTCT AGCATGA

5 **Figure 5D DNA and amino acid sequence list****SEQ. ID. NO. 2** Amino acid sequence of DNA polymerase for HSV2-MS-M1

1 MFCAAGGPTS PGGKSAARAA SGFFAPHNPR GATQTAPPPC RRQNFYNPHL
 10 51 AQTGTQPKAP GPAQRHTYYS ECDEFRIAP RSLDEDAPAE QRTGVHDGRL
 101 RRAPKVYCGG DERDVLRVGP EGFWRRLRL WGGADHAPKG FDPTVTVFHV
 15 151 YDILEHVEHA YSMRAAQLHE RFMDAITPAG TVITLLGLTP EGHRAVAVHV
 201 GTRQYFYMNK AEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE
 251 AEVVERADVY YYETRPTLYY RVFVRSGRAL AYLCDNFCPA IRKYEGGVDA
 20 301 TTRFILDNPG FVTFGWYRLK PGRGNAPAQ RPPTAFGTSS DVEFNCTADN
 351 LAVEGAMCDL PAYKLMCFDI ECKAGGEDEL AFPVAERPED LVIQISCLLY
 401 DLSTTALEHI LLFSLGSCDL PESHLSDLAS RGLPAPVVLE FDSEFEMLLA
 25 451 FMTFVKQYGP EFVTGYNIIN FDWPFVLTCL TEIYKVPLDG YGRMNRRGVF
 501 RVWDIGQSHF QKRSEIKVNG MVNIDMYGII TDKVKLSSYK LNAVAEAVLK
 30 551 DKKKDLSDYRD IPAYYASGPA QRGVIGEYCV QDSLLVGQLF FKFLPHLELS
 601 AVARLAGINI TRTIYDGQOI RVFTCLLRLA GQKGFILPDT QGRFRGLDKE
 651 APKRPVPRG EGERPGDGNG DEDKDDDEDE DGDEREEVAR ETGGRHVGYQ
 35 701 GARVLDPTSG FHVDPVVVFD FASLYPSIIQ AHNLCFSTLS LRPEAVAHLE
 751 ADRDYLEIEV GGRRLFFVKA HVRESLLSIL LRDWLAMRKQ IRSRIPQSTP
 40 801 EEAVLLDKQQ AAIKVCNSV YGFTGAQHGL LPCLHVAATV TTIGREMLLA
 851 TRAYVHARWA EFDQLLADFP EAAGMRAPGP YSMRIYGDT DSIFVLCRGL
 901 TAAGLVAMGD KMASHISRAL FLPIKLECE KTFTKLLLIA KKKYIGVICG
 45 951 GKMLIKGVDL VRKNNCAFIN RTSRALVDLL FYDDTVSGAA AALAERPAEE
 1001 WLARPLPEGL QAFGAVLVDA HRRITDPERD IQDFVLTAEI SRHPRAYTNK

 50 1051 RLAHLTVYYK LMARRAQVPS IKDRIPYVIV AQTREVEETV ARLAALRELD
 1101 AAAPGDEPAP PAALPSPAKR PRETPSHADP PGGASKPRKL LVSELAEDPG
 1151 YAIARGVPLN TDYYFSHLLG AACVTFKALF GNNAKITESL LKRFIPETWH
 55 1201 PPDDVAARLR AAGFGPAGAG ATAEETRRML HRAFDTLA*

5 **Figure 5E DNA and amino acid sequence list****SEQ.ID.NO. 3** DNA sequence of DNA polymerase gene for HSV2-186-M1

10 1 ATGTTTTGTG CCGCGGGCGG CCGGGCTTCC CCCGGGGGGA AGTCGGCGGC
 51 TCGGGCGGCG TCTGGGTTTT TTGCCCCCA CAACCCCGG GGAGCCACCC
 101 AGACGGCACC GCCGCCTTGC CGCGGCAGA ACTTCTACAA CCCCACCTC
 15 151 GCTCAGACCG GAACGCAGCC AAAGGCCCCC GGGCCGGCTC AGCGCCATAC
 201 GTACTACAGC GAGTGCACG AATTTCGATT TATCGCCCCG CGTTCGCTGG
 20 251 ACGAGGACGC CCCCGCGGAG CAGCGCACCG GGGTCCACGA CGGCCGCTC
 301 CGGCGCGCCC CTAAGGTGTA CTGCGGGGGG GACGAGCGCG ACGTCCTCCG
 351 CGTGGGCCCC GAGGGCTTCT GGCCGCGTCG CTTGCGCTG TGGGGCGGTG
 25 401 CGGACCATGC CCCCAGGGG TCGACCCA CCGTCACCGT CTTCCACGTG
 451 TACGACATCC TGGAGCACGT GGAACACGCG TACAGCATGC GCGCCGCCA
 501 GCTCCACGAG CGATTTATGG ACGCCATCAC GCCCGCCGGG ACCGTCATCA
 551 CGCTTCTGGG TCTGACCCCC GAAGGCCATC GCGTCGCCGT TCACGTCTAC
 601 GGCACGCGGC AGTACTTTTA CATGAACAAG GCGGAGGTGG ATCGGCACCT
 35 651 GCAGTGCCGT GCCCCGCGCG ATCTCTGCGA GCGCCTGGCG GCGGCCCTGC
 701 GCGAGTCGCC GGGGGCGTCG TTCCGCGGCA TCTCCGCGGA CCACTTCGAG
 40 751 GCGGAGGTGG TGGAGCGCGC CGACGTGTAC TATTACGAAA CGCGCCCGAC
 801 CCTGTACTAC CGCGTCTTCG TGCGAAGCGG GCGCGCGCTG GCCTACCTGT
 851 GCGACAACTT TTGCCCCGCG ATCAGGAAGT ACGAGGGGGG CGTCGACGCC
 45 901 ACCACCCGGT TTATCCTGGA CAACCCGGGG TTTGTACCT TCGGCTGGTA
 951 CCGCCTCAAG CCCGGCCGCG GGAACGCGCC GGCCCAACCG CGCCCCCGA
 50 1001 CGGCGTTCGG AACCTCGAGC GACGTCGAGT TAACTGCAC GGCGGACAAC
 1051 CTGGCCGTCG AGGGGGCCAT GTGTGACCTG CCGGCCTACA AGCTCATGTG
 1101 CTTGATATC GAATGCAAGG CCGGGGGGGA GGACGAGCTG GCCTTTCCGG
 55 1151 TCGCGGAACG CCCGGAAGAC CTCGTCATCC AGATCTCCTG TCTGCTCTAC
 1201 GACCTGTCCA CCACCGCCT CGAGCACATC CTCCTGTTTT CGCTCGGATC

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Figure 5F DNA and amino acid sequence list

10 1251 CTGCGACCTC CCGAGTCCC ACCTCAGCGA TCTCGCCTCC AGGGGCCTGC
 1301 CGGCCCCCGT CGTCCTGGAG TTTGACAGCG AATTCGAGAT GCTGCTGGCC
 1351 TTCATGACCT TCGTCAAGCA GTACGGCCCC GAGTTCGTGA CCGGGTACAA
 15 1401 CATCATCAAC TTCGACTGGC CCTTCGTCCT GACCAAGCTG ACGGAGATCT
 1451 ACAAGGTCCC GCTCGACGGG TACGGGCGCA TGAACGGCCG GGGTGTGTTC
 20 1501 CGCGTGTGGG ACATCGGCCA GAGCCACTTT CAGAAGCGCA GCAAGATCAA
 1551 GGTGAACGGG ATGGTGAACA TCGACATGTA CGGCATCATC ACGACAAGG
 1601 TCAAACCTCTC CAGCTACAAG CTGAACGCCG TCGCCGAGGC CGTCTTGAAG
 25 1651 GACAAGAAGA AGGATCTGAG CTACCGCGAC ATCCCCGCCT ACTACGCCTC
 1701 CGGGCCCGCG CAGCGCGGGG TGATCGGCGA GTATTGTGTG CAGGACTCGC
 30 1751 TGCTGGTCGG GCAGCTGTTC TTCAAGTTTC TGCCGCACCT GGAGCTTTCC
 1801 GCCGTCGCGC GCCTGGCGGG CATCAACATC ACCCGCACCA TCTACGACGG
 1851 CCAGCAGATC CGCGTCTTCA CGTGCCTCCT GCGCCTTGCG GGCCAGAAGG
 35 1901 GCTTCATCCT GCCGGACACC CAGGGGCGGT TTCGGGGCCT CGACAAGGAG
 1951 GCGCCCAAGC GCCCGGCCGT GCCTCGGGGG GAAGGGGAGC GGCCGGGGGA
 40 2001 CGGGAACGGG GACGAGGATA AGGACGACGA CGAGGACGGG GACGAGGACG
 2051 GGGACGAGCG CGAGGAGGTC GCGCGCGAGA CCGGGGGCCG GCACGTTGGG
 2101 TACCAGGGGG CCCGGGTCCT CGACCCACACC TCCGGGTTTC ACGTCGACCC
 45 2151 CGTGGTGGTG TTTGACTTTG CCAGCCTGTA CCCCAGCATC ATCCAGGCCC
 2201 ACAACCTGTG CTTAGTACG CTCTCCCTGC GGCCCGAGGC CGTCGCGCAC
 50 2251 CTGGAGGCGG ACGGGACTA CCTGGAGATC GAGGTGGGGG GCCGACGGCT
 2301 GTTCTTCGTG AAGGCCACG TACGCGAGAG CCTGCTGAGC ATCCTGCTGC
 2351 GCGACTGGCT GGCCATGCGA AAGCAGATCC GCTCGCGGAT CCCCCAGAGC
 55 2401 CCCCCGAGG AGGCCGTCCT CCTCGACAAG CAACAGGCCG CCATCAAGGT
 2451 GGTGTGCAAC TCGGTGTACG GGTTCACCGG GGCGCAGCAC GGTCTTCTGC
 60 2501 CCTGCCTGCA CGTGGCCGCC ACCGTGACGA CCATCGGCCG CGAGATGCTC

5 **Figure 5G DNA and amino acid sequence list**

2551 CTCGCGACGC GCGCGTACGT GCACGCGCGC TGGGCGGAGT TCGATCAGCT
 10 2601 GCTGGCCGAC TTTCCGGAGG CGGCCGGCAT GCGCGCCCCC GGTCCGTACT
 2651 CCATGCGCAT CATCTACGGG GACACGGACT CCATTTTCGT TTTGTGCCGC
 2701 GGCCTCACGG CCGCGGGCCT GGTGGCCATG GGCGACAAGA TGGCGAGCCA
 15 2751 CATCTCGCGC GCGCTGTTCC TCCCCCGAT CAAGCTCGAG TGCGAAAAAA
 2801 CGTTCACCAA GCTGCTGCTC ATCGCCAAGA AAAAGTACAT CGGCGTCATC
 20 2851 TGCGGGGGCA AGATGCTCAT CAAGGGCGTG GATCTGGTGC GCAAAAACAA
 2901 CTGCGCGTTT ATCAACGCA CCTCCAGGGC CCTGGTCGAC CTGCTGTTTT
 2951 ACGACGATAC CGTATCCGGA GCGGCCGCG CGTTAGCCGA GCGCCCCGCA
 25 3001 GAGGAGTGGC TGGCGCGACC CCTGCCCCGAG GGAAGTGCAGG CGTTCGGGGC
 3051 CGTCCTCGTA GACGCCCATC GGCGCATCAC CGACCCGAG AGGGACATCC
 30 3101 AGGACTTTGT CCTCACCGCC GAACTGAGCA GACACCCGCG CGCGTACACC
 3151 AACAAGCGCC TGGCCCACCT GACGGTGTAT TACAAGCTCA TGGCCCCGCG
 3201 CGCGCAGGTC CCGTCCATCA AGGACCGGAT CCCGTACGTG ATCGTGGCCC
 35 3251 AGACCCGCGA GGTAGAGGAG ACGGTCGCGC GGCTGGCCGC CCTCCGCGAG
 3301 CTAGACGCGG CCGCCCCAGG GGACGAGCCG GCGCCCCCAG CGGCCCTGCC
 40 3351 CTCCCCGGCC AAGCGCCCCG GGGAGACGCC GTGCGATGCC GACCCCCCGG
 3401 GAGGCGCGTC CAAGCCCCGC AAGCTGCTGG TGTCCGAGCT GGCGGAGGAT
 3451 CCCGGGTACG CCATCGCCCC GGGCGTTCCG CTCAACACGG ACTATTACTT
 45 3501 CTCGCACCTG CTGGGGGCGG CCTGCGTGAC GTTCAAGGCC CTGTTTGGAA
 3551 ATAACGCCAA GATCACCGAG AGTCTGTAA AGAGGTTTAT TCCCGAGACG
 50 3601 TGGCACCCCC CGGACGACGT GGCCGCGCGG CTCAGGGCCG CGGGGTTTCG
 3651 GCCGGCGGGG GCCGGCGCTA CGGCGGAGGA AACTCGTCGA ATGTTGCATA
 3701 GAGCCTTTGA TACTCTAGCA TGA
 55

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Figure 5H DNA and amino acid sequence list**SEQ.ID.NO. 4 Amino acid sequence of DNA polymerase for HSV2-186-M1**

10 1 MFCAAGGPAS PGGKSAARAA SGFFAPHNPR GATQTAPPPC RRQNFYNPHL
 51 AQTGTQPKAP GPAQRHTYYS ECDEFRIAP RSLDEDAPAE QRTGVHDGRL
 101 RRAPKVYCGG DERDVLRVGP EGFWRRLRL WGGADHAPEG FDPTVTVFHV
 15 151 YDILEHVEHA YSMRAAQLHE RFMDAITPAG TVITLLGLTP EGHRAVHVY
 201 GTRQYFYMNK AEVDRHLQCR APRDLCERLA AALRESPGAS FRGISADHFE
 20 251 AEVVERADVY YYETRPTLYY RVFVRSGRAL AYLCDNFCA IRKYEGGVDA
 301 TTRFILDNPG FVTFGWYRLK PGRGNAPAQ RPPTAFGTSS DVEFNCTADN
 351 LAVEGAMCDL PAYKLMCFDI ECKAGGEDEL AFPVAERPED LVIQISCLLY
 25 401 DLSTTALEHI LLFSLGSCDL PESHLSDLAS RGLPAPVVLE FDSEFEMLLA
 451 FMTFVKQYGP EFVTGYNIIN FDWPFVLTKL TEIYKVPLDG YGRMNNGRGVF
 30 501 RVWDIGQSHF QKRSEIKVNG MVNIDMYGII TDKVKLSSYK LNAVAEAVLK
 551 DKKKDLSDYR IPAYYASGPA QRGVIGEYCV QDSLLVGQLF FKFLPHLELS
 601 AVARLAGINI TRTIYDGQOI RVFTCLRLA GQKGFILPDT QGRFRGLDKE
 35 651 APKRPAVPRG EGERPGDGNG DEDKDDDEDG DEDGDREEV ARETGGRHVG
 701 YQGARVLDPT SGFHVDPVVV FDFASLYPSI IQAHNLCFST LSLRPEAVAH
 40 751 LEADRDYLEI EVGGRRLLFFV KAHVRESLLS ILLRDWLAMR KQIRSRIPOS
 801 PPEEAVLLDK QQAAIKVVCN SVYGFTGAQH GLLPCLHVAA TVTTIGREML
 851 LATRAYVHAR WAEFDQLLAD FPEAAGMRAP GPYSMRITYG DTDSIFVLCR
 45 901 GLTAAGLVAM GDKMASHISR ALFLPPIKLE CEKTFTKLLL IAKKKYIGVI
 951 CGGKMLIKGV DLVRKNNCAF INRTSRALVD LLFYDDTVSG AAAALAERPA

 50 1001 EEWLARPLPE GLQAFGAVLV DAHRRITDPE RDIQDFVLTA ELSRHPRAYT
 1051 NKRLAHLTVY YKLMARRAQV PSIKDRIPYV IVAQTREVEE TVARLAALRE
 1101 LDAAAPGDEP APPAALPSA KRPRETPSHA DPPGGASKPR KLLVSELAED
 55 1151 PGYAIARGVP LNTDYYFSLH LGAACVTFKA LFGNNAKITE SLLKRFIPET
 1201 WHPPDDVAAR LRAAGFGPAG AGATAEETRR MLHRAFDTLA *

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5 **Figure 5I DNA and amino acid sequence list**

SEQ.ID.NO. 5 DNA sequence of DNA polymerase gene for HSV1-KOS-M1

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10   1 ATGTTTTC CG GTGGCGGCGG CCGCTGTCC CCGGAGGAA AGTCGGCGGC
    51 CAGGGCGGCG TCGGGTTTT TTGCGCCCGC CGGCCCTCGC GGAGCCGGCC
    101 GGGGACCCCC GCCTTGTTTG AGGCAAACT TTTACAACCC CTACCTCGCC
15   151 CCAGTCGGGA CGCAACAGAA GCGACCGGG CCAACCCAGC GCCATACGTA
    201 CTATAGCGAA TCGATGAAT TCGATT CAT CGCCCCGCGG GTGCTGGACG
    251 AGGATGCCCC CCGGAGAAG CGCGCCGGG TGCACGACGG TCACCTCAAG
    301 CGCGCCCCCA AGGTGTACTG CGGGGGGGAC GAGCGCGACG TCCTCCGCGT
    351 CGGGTCGGGC GGCTTCTGGC CGCGGCGCTC GCGCCTGTGG GCGGCGTGG
25   401 ACCACGCCCC GCGGGGGTTC AACCCACCG TCACCGTCTT TCACGTGTAC
    451 GACATCCTGG AGAACGTGGA GCACGCGTAC GGCATGCGCG CGGCCAGTT
    501 CCACGCGCGG TTTATGGACG CCATCACACC GACGGGGACC GTCATCACGC
    551 TCCTGGGCCT GACTCCGGA GGCCACCGG TGGCCGTTCA CGTTTACGGC
    601 ACGCGGCAGT ACTTTTACAT GAACAAGGAG GAGGTTGACA GGCACCTACA
35   651 ATGCCGCGCC CCACGAGATC TCTGCGAGCG CATGGCCGCG GCCCTGCGCG
    701 AGTCCCCGGG CGCGTCGTT CGCGGCATCT CCGCGGACCA CTTCGAGGCG
    751 GAGGTGGTGG AGCGCACCGA CGTGTACTAC TACGAGACGC GCCCCGCTCT
    801 GTTTTACCGC GTCTACGTCC GAAGCGGGCG CGTGCTGTG TACCTGTGCG
    851 ACAACTTCTG CCCGGCCATC AAGAAGTACG AGGGTGGGGT CGACGCCACC
45   901 ACCCGGTTCA TCCTGGACAA CCCCGGGTTC GTCACCTTCG GCTGGTACCG
    951 TCTCAAACCG GGCCGGAACA ACACGCTAGC CCAGCCGCGG GCCCCGATGG
    1001 CCTTCGGGAC ATCCAGCGAC GTCGAGTTTA ACTGTACGGC GGACAACCTG
    1051 GCCATCGAGG GGGGCATGAG CGACCTACCG GCATACAAGC TCATGTGCTT
    1101 CGATATCGAA TGCAAGGCGG GGGGGGAGGA CGAGCTGGCC TTTCGGGTGG
55   1151 CCGGGCACCC GGAGGACCTG GTTATT CAGA TATCCTGTCT GCTCTACGAC
    1201 CTGTCCACCA CCGCCCTGGA GCACGTCCTC CTGTTTTCGC TCGGTTCTG

```

5

Figure 5J DNA and amino acid sequence list

10 1251 CGACCTCCCC GAATCCCACC TGAACGAGCT GGCGGCCAGG GGCCTGCCCCA
 1301 CGCCCGTGGT TCTGGAATTC GACAGCGAAT TCGAGATGCT GTTGGCCTTC
 1351 ATGACCCTTG TGAAACAGTA CGGCCCCGAG TTCGTGACCG GGTACAACAT
 15 1401 CATCAACTTC GACTGGCCCT TCTTGCTGGC CAAGTTGACG GACATTTACA
 1451 AGGTCCCCCT GGACGGGTAC GGCCGCATGA ACGGCCGGGG CGTGTTTCGC
 20 1501 GTGTGGGACA TAGGCCAGAG CCACTTCCAG AAGCGCAGCA AGATAAAGGT
 1551 GAACGGCATG GTGAACATCG ACATGTACGG GATCATAACC GACAAGATCA
 1601 AGCTCTCGAG CTACAAGCTC AACGCCGTGG CGAAGCOGT CCTGAAGGAC
 25 1651 AAGAAGAAGG ACCTGAGCTA TCGCGACATC CCCGCCTACT ACGCCGCCGG
 1701 GCCCGCGCAA CGCGGGGTGA TCGGCGAGTA CTGCATACAG GATTCCCTGC
 30 1751 TGGTGGGCCA GCTGTTTTTT AAGTTTTTGC CCCATCTGGA GCTCTCGGCC
 1801 GTCGCGCGCT TGGCGGGTAT TAACATCACC CGCACCATCT ACGACGGCCA
 1851 GCAGATCCGC GTCTTTACGT GCCTGCTGCG CCTGGCCGAC CAGAAGGGCT
 35 1901 TTATTCTGCC GGACACCCAG GGGCGATTTA GGGGCGCCGG GGGGGAGGCG
 1951 CCCAAGCGTC CGGCCGAGC CCGGGAGGAC GAGGAGCGGC CAGAGGAGGA
 40 2001 GGGGGAGGAC GAGGACGAAC GCGAGGAGGG CGGGGGCGAG CGGGAGCCGG
 2051 AGGGCGCGCG GGAGACCGCC GGCCGGCACG TGGGGTACCA GGGGGCCAGG
 2101 GTCCTTGACC CCACTTCCGG GTTTCACGTG AACCCCGTGG TGGTGTTCGA
 45 2151 CTTTGCCAGC CTGTACCCCA GCATCATCCA GGCCCAACAAC CTGTGCTTCA
 -----2201 GCACGCTETC CCTGAGGGCC GACGCAGTGG CGCACCTGGA GGCGGGCAAG-----
 50 2251 GACTACCTGG AGATCGAGGT GGGGGGGCGA CGGCTGTTCT TCGTCAAGGC
 2301 TCACGTGCGA GAGAGCCTCC TCAGCATCCT CCTGCGGGAC TGGCTCGCCA
 2351 TGCGAAAGCA GATCCGCTCG CGGATTCCCC AGAGCAGCCC CGAGGAGGCC
 55 2401 GTGCTCCTGG ACAAGCAGCA GGCCGCCATC AAGGTCGTGT GTAACCTCGGT
 2451 GTACGGGTTC ACGGGAGCGC AGCACGGA CTGCGCGTGC CTGCACGTTG
 60 2501 CCGCGACGGT GACGACCATC GGCCGCGAGA TGCTGCTCGC GACCCGCGAG

5 **Figure 5K DNA and amino acid sequence list**

2551 TACGTCCACG CGCGCTGGGC GGCCTTCGAA CAGCTCCTGG CCGATTTCCT
 10 2601 GGAGGCGGCC GACATGCGCG CCCC CGGGCC CTATTCCATG CGCATCATCT
 2651 ACGGGGACAC GGACTCCATA TTTGTGCTGT GCCGCGGCCT CACGGCCGCC
 2701 GGGCTGACGG CCATGGGCGA CAAGATGGCG AGCCACATCT CGCGCGCGCT
 15 2751 GTTTCTGCCC CCCATCAAAC TCGAGTGCGA AAAGACGTTC ACCAAGCTGC
 2801 TGCTGATCGC CAAGAAAAAG TACATCGGCG TCATCTACGG GGGTAAGATG
 20 2851 CTCATCAAGG GCGTGGATCT GGTGCGCAAA AACAACTGCG CGTTTATCAA
 2901 CCGCACCTCC AGGGCCCTGG TCGACCTGCT GTTTTACGAC GATACCGTAT
 2951 CCGGAGCGGC CGCCGCGTTA GCCGAGCGCC CCGCAGAGGA GTGGCTGGCG
 25 3001 CGACCCCTGC CCGAGGGACT GCAGGCGTTC GGGGCCGTCC TCGTAGACGC
 3051 CCATCGGCGC ATCACCGACC CGGAGAGGGA CATCCAGGAC TTTGTCCTCA
 30 3101 CCGCCGA ACT GAGCAGACAC CGCGCGCGT ACACCAACAA GCGCCTGGCC
 3151 CACCTGACGG TGTATTACAA GTCATGGCC CGCCGCGCGC AGGTCCCGTC
 3201 CATCAAGGAC CGGATCCCGT ACGTGATCGT GGCCAGACC CGCGAGGTAG
 35 3251 AGGAGACGGT CGCGCGGCTG GCCGCCCTCC GCGAGCTAGA CGCCGCGGCC
 3301 CCAGGGGACG AGCCCGCCCC CCCC GCGGCC CTGCCCTCCC CGGCCAAGCG
 40 3351 CCCCCGGGAG ACGCCGTCGC ATGCCGACCC CCCGGGAGGC GCGTCCAAGC
 3401 CCGCAAGCT GCTGGTGTCC GAGCTGGCCG AGGATCCCGC ATACGCCATT
 3451 GCCACGGCG TCGCCCTGAA CACGGACTAT TACTTCTCCC ACCTGTTGGG
 45 3501 GCGGCGTGC GTGACATTCA AGGCCCTGTT TGGGAATAAC GCCAAGATCA
 3551 CCGAGAGTCT GTTAAAAAGG TTTATTCCCG AAGTGTGGCA CCCCCGGAC
 50 3601 GACGTGGCCG CGCGGCTCCG GGCCG CAGGG TTCGGGGCGG TGGGTGCCG
 3651 CGCTACGGCG GAGGAACTC GTCGAATGTT GCATAGAGCC TTTGATACTC
 3701 TAGCATGA
 55
 60

5 **Figure 5L DNA and amino acid sequence list****SEQ.ID.NO. 6** Amino acid sequence of DNA polymerase for HSV1-KOS-M1

1 MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGRGPPPCL RQNFYNPYLA
 10 51 PVGTQQKPTG PTQRHTYYSE CDEFRIAPR VLDEDAPEK RAGVHDGHLK
 101 RAPKVYCGGD ERDVLRVGSG GFWPRRSRLW GGVDHAPAGF NPTVTVFHVY
 15 151 DILENVEHAY GMRAAQFHAR FMDAITPTGT VITLLGLTPE GHRVAVHVYG
 201 TRQYFYMNKE EVDRHLQCRA PRDLCERMAA ALRESPGASF RGISADHFEA
 251 EVVERTDVYY YETRPALFYR VYVRSGRVLS YLCDNFCPAI KKYEGGV DAT
 20 301 TRFILDNPGF VTFGWYRLKP GRNNTLAQPR APMAFGTSSD VEFNCTADNL
 351 AIEGGMSDLP AYKLMCFDIE CKAGGEDELA FPVAGHPEDL VIQISCLLYD
 401 LSTTALEHVL LFSLGSCDLP ESHLNELAAR GLPTPVVLEF DSEFEMLLAF
 25 451 MTLVKQYGPE FVTGYNIINF DWPFLAKLT DIYKVPLDGY GRMN GRGVFR
 501 VWDIGQSHFQ KRSKIKVNGM VNIDMYGIIT DKIKLSSYKL NAVA EAVLKD
 30 551 KKKDLSYRDI PAYYAAGPAQ RGVIGEYCIQ DSLLVGQLFF KFLPHLELSA
 601 VARLAGINIT RTIYDGQQIR VFTCLRLAD QKGFILPDTQ GRFRGAGGEA
 651 PKRPAAARED EERPEEEGED EDEREEGGGE REPEGARETA GRHVGYQGAR
 35 701 VLDPTSGFHV NPVVVFDFAS LYPSTIAHN LCFSTLSLRA DAVAHLEAGK
 751 DYLEIEVGGR RLFFVKAHVR ESLLSILLRD WLAMRKQIRS RIPQSSPEEA
 40 801 VLLDKQQA AI KVCNSVYGF TGAQHGLLPC LHVAATVTTI GREMLLATRE
 851 YVHARWAAFE QLLADFPEAA DMRAPGPYSM RIYGD TDSI FVLCRGLTAA
 901 GLTAMGDKMA SHISRALFLP PIKLECEKTF TKLLLIAKKK YIGVITYGGKM
 45 951 LIKGVDLVRK NNCAFINRTS RALVDLLFYD DTVSGAAAAL AERP AEWLA
 1001 RPLPEGLQAF GAVLVDAHRR ITDPERDIQD FVLTAELSRH PRAYTNKRLA
 50 1051 HLTVYYKLMA RRAQVPSIKD RPYVIVAQT REVEETVARL AALRELDAAA
 1101 PGDEPAPPAA LPSPAKRPRE TPSHADPPGG ASKPRKLLVS ELAEDPAYAI
 1151 AHGVALNTDY YFSHLLGAAC VTFKALFGNN AKITESLLKR FIPEVWHPPD
 55 1201 DVAARLRAAG FGAVGAGATA EETRRMLHRA FDTLA*

5 **Figure 5M DNA and amin acid sequence list****SEQ.ID.NO. 7** DNA sequence of HSV polymerase gene for HSV1-F-M1

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1  ATGTTTTCCG GTGGCGGCGG CCCGCTGTCC CCCGAGGAA AGTCGGCGGC
10  51  CAGGGCGGCG TCCGGGTTTT TTGCGCCCGC CGGCCCTCGC GGAGCCGGCC
    101  GGGGACCCCC GCCTTGCTTG AGGCAAACT TTTACAACCC CTACCTCGCC
15  151  CCAGTCGGA CGAACAGAA GCCGACCGG CCAACCCAGC GCCATACGTA
    201  CTATAGCGAA TCGATGAAT TTCGATTCAT CGCCCCGCGG GTGCTGGACG
    251  AGGATGCCCC CCCGAGAAG CGCGCCGGG TGCACGACG TCACCTCAAG
20  301  CGCGCCCCCA AGGTGTACTG CGGGGGGAC GAGCGCGAC TCCTCCGCGT
    351  CGGTCGGGC GGCTTCTGGC CGCGCGCTC GCGCTGTGG GCGGCGGTGG
25  401  ACCACGCCCC GCGGGGTTC AACCCACCG TCACCGTCTT TCACGTGTAC
    451  GACATCCTGG AGAACGTGA GCACGCGTAC GGCATGCGCG CGGCCAGTT
    501  CCACGCGCGG TTTATGGACG CCATCACACC GACGGGGACC GTCATCACGC
30  551  TCCTGGGCT GACTCCGAA GGCCACCGG TGGCCGTTC CGTTTACGGC
    601  ACGCGCAGT ACTTTTACAT GAACAAGGAG GAGGTCGACA GGCACCTACA
35  651  ATGCCGCGCC CCACGAGATC TCTGCGAGCG CATGGCCGCG GCCCTGCGCG
    701  AGTCCCCGGG CGCGTCGTTT CGCGGCATTT CCGCGGACCA CTTGAGGGC
    751  GAGGTGGTGG AGCGACCGA CGTGCTACTAC TACGAGACGC GCCCCGCTCT
40  801  GTTTTACCGC GTCTACGTCC GAAGCGGGCG CGTGCTGTCG TACCTGTGCG
    851  ACAACTTCTG CCCGGCCATC AAGAAGTACG AGGTGGGGT CGACGCCACC
45  901  ACCCGTTCA TCCTGGACAA CCCCGGGTTT GTCACCTTCG GCTGGTACCG
    951  TCTCAAACCG GGCCGAACA ACACGCTAGC CCAGCCGCGG GCCCCGATGG
1001 CCTTCGGGAC ATCCAGCGAC GTCGAGTTTA ACTGTACGGC GGACAACCTG
50  1051 GCCATCGAGG GGGGCATGAG CGACCTACCG GCATACAAGC TCATGTGCTT
    1101 CGATATCGAA TGCAAGGCGG GGGGGAGGA CGAGCTGGCC TTCCGGTGG
55  1151 CCGGGCACCC GGAGGACCTG GTCATCCAGA TATCCTGTCT GCTCTACGAC
    1201 CTGTCCACCA CCGCCCTGGA GCACGTCCTC CTGTTTTCGC TCGGTTCTTG
    1251 CGACCTCCCC GAATCCCACC TGAACGAGCT GGCGCCAGG GGCCTGCCCA
60

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5 **Figure 5N DNA and amino acid sequence list**

| | | | | | | |
|----|------|-------------|-------------|------------|------------|------------|
| | 1301 | CGCCCGTGGT | TCTGGAATTC | GACAGCGAAT | TCGAGATGCT | GTTGGCCTTC |
| | 1351 | ATGACCCCTTG | TGAAACAGTA | CGGCCCCGAG | TTCGTGACCG | GGTACAACAT |
| 10 | 1401 | CATCAACTTC | GACTGGCCCT | TCTTGCTGGC | CAAGCTGACG | GACATTTACA |
| | 1451 | AGGTCCCCCT | GGACGGGTAC | GGCCGCATGA | ACGGCCGGGG | CGTGTTCGCG |
| 15 | 1501 | GTGTGGGACA | TAGGCCAGAG | CCACTTCCAG | AAGCGCAGCA | AGATAAAGGT |
| | 1551 | GAACGGCATG | GTGAACATCG | ACATGTACGG | GATTATAACC | GACAAGATCA |
| | 1601 | AGCTCTCGAG | CTACAAGCTC | AACGCCGTGG | CCGAAGCCGT | CCTGAAGGAC |
| 20 | 1651 | AAGAAGAAGG | ACCTGAGCTA | TCGCGACATC | CCCGCCTACT | ACGCCGCCGG |
| | 1701 | GCCCCGCGAA | CGCGGGGTGA | TCGGCGAGTA | CTGCATACAG | GATTCCCTGC |
| 25 | 1751 | TGGTGGGCCA | GCTGTTTTTT | AAGTTTTTGC | CCCATCTGGA | GCTCTCGGCC |
| | 1801 | GTCGCGCGCT | TGGCGGGTAT | TAACATCACC | CGCACCATCT | ACGACGGCCA |
| | 1851 | GCAGATCCGC | GTCTTTACGT | GCCTGCTGCG | CCTGGCCGAC | CAGAAGGGCT |
| 30 | 1901 | TTATTCTGCC | GGACACCCAG | GGGCGATTTA | GGGGCGGCGG | GGGGGAGGCG |
| | 1951 | CCCAAGCGTC | CGGCCCGCAGC | CCGGGAGGAC | GAGGAGCGGC | CAGAGGAGGA |
| 35 | 2001 | GGGGGAGGAC | GAGGACGAAC | GCGAGGAGGG | CGGGGGCGAG | CGGGAGCCGG |
| | 2051 | AGGGCGCGCG | GGAGACCGCC | GGCCGGCACG | TGGGGTACCA | GGGGGCCAGG |
| | 2101 | GTCCTTGACC | CCACTTCCGG | GTTTCATGTG | AACCCCGTGG | TGGTGTTCGA |
| 40 | 2151 | CTTTGCCAGC | CTGTACCCCA | GCATCATCCA | GGCCCACAAC | CTGTGCTTCA |
| | 2201 | GCACGCTCTC | CCTGAGGGCC | GACGCAGTGG | CGCACCTGGA | GGCGGGCAAG |
| 45 | 2251 | GACTACCTGG | AGATCGAGGT | GGGGGGGCGA | CGGCTGTTCT | TCGTCAAGGC |
| | 2301 | TCACGTGCGA | GAGAGCCTCC | TCAGCATCCT | CCTGCGGGAC | TGGCTCGCCA |
| | 2351 | TGCGAAAGCA | GATCCGCTCG | CGGATTCCCC | AGAGCAGCCC | CGAGGAGGCC |
| 50 | 2401 | GTGCTCCTGG | ACAAGCAGCA | GGCCGCCATC | AAGGTCGTGT | GTAACTCGGT |
| | 2451 | TTACGGGTTC | ACGGGAGCGC | AGCACGGACT | CCTGCCGTGC | CTGCACGTTG |
| 55 | 2501 | CCGCGACGGT | GACGACCATC | GGCCGCGAGA | TGCTGCTCGC | GACCCGCGAG |
| | 2551 | TACGTCCACG | CGCGCTGGGC | GGCCTTCGAA | CAGCTCCTGG | CCGATTTCCC |
| 60 | 2601 | GGAGGCGGCC | GACATGCGCG | CCCCGGGGCC | CTATTCCATG | CGCATCATCT |

5

Figure 50 DNA and amino acid sequence list

10 2651 ACGGGGACAC GGACTCCATC TTTGTGCTGT GCCGCGGCCT CACGGCCGCC
 2701 GGGCTGACGG CCGTGGGCGA CAAGATGGCG AGCCACATCT CGCGCGCGCT
 2751 GTTCTGTGCC CCCATCAAAC TCGAGTGC GA AAAGACGTTC ACCAAGCTGC
 15 2801 TGCTGATCGC CAAGAAAAAG TACATCGGCG TCATCTACGG GGGTAAGATG
 2851 CTCATCAAGG GCGTGGATCT GGTGCGCAAA AACAACTGCG CGTTTATCAA
 2901 CCGCACCTCC AGGGCCCTGG TCGACCTGCT GTTTTACGAC GATACCGTAT
 20 2951 CCGGAGCGGC CGCCGCGTTA GCCGAGCGCC CCGCAGAGGA GTGGCTGGCG
 3001 CGACCCCTGC CCGAGGGACT GCAGGCGTTC GGGGCCGTCC TCGTAGACGC
 25 3051 CCATCGGCGC ATCACCAGAC CGGAGAGGGA CATCCAGGAC TTTGTCCTCA
 3101 CCGCCGAAC T GAGCAGACAC CCGCGCGCGT ACACCAACAA GCGCCTGGCC
 3151 CACCTGACGG TGTATTACAA GTCATGGCC CGCCGCGCGC AGGTCCCGTC
 30 3201 CATCAAGGAC CGGATCCCGT ACGTGATCGT GGCCCAGACC CGCGAGGTAG
 3251 AGGAGACGGT CGCGCGGCTG GCCGCCCTCC GCGAGCTCGA CGCCGCCGCC
 35 3301 CCAGGGGACG AGCCCGCCCC CCCCGCGGCC CTGCCCTCCC CGGCCAAGCG
 3351 CCCCCGGGAG ACGCCGTTGC ATGCCGACCC CCCGGGAGGC GCGTCCAAGC
 3401 CCCGCAAGCT GCTGGTGTCC GAGCTGGCCG AGGATCCCGC ATACGCCATT
 40 3451 GCCCACGGCG TCGCCCTGAA CACGGACTAT TACTTCTCCC ACCTGTTGGG
 3501 GGCGGCGTGC GTGACATTCA AGGCCCTGTT TGGGAATAAC GCCAAGATCA
 45 3551 CCGAGAGTCT GTTAAAAAGG TTTATTCCCG AAGTGTGGCA CCCCCCGGAC
 3601 GACGTGGCCG CGCGGCTCCG GGCCGCAGGG TTCGGGGCGG TGGGTGCCGG

 50 3651 CGCTACGGCG GAGGAACTC GTCGAATGTT GCATAGAGCC TTTGATACTC
 3701 TAGCATGA

5 **Figure 5P DNA and amino acid sequence list**

SEQ.ID.NO. 8 Amino acid sequence of DNA polymerase for HSV1-F-M1

10 1 MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGRGPPPCL RQNFYNPYLA
 51 PVGTQQKPTG PTQRHTYYSE CDEFRFIAPR VLDEDAPEEK RAGVHDGHLK
 101 RAPKVYCGGD ERDVLRVGSG GFWPRRSRLW GGVDHAPAGF NPTVTVFHVY
 15 151 DILENVEHAY GMRAAQFHAR FMDAITPTGT VITLLGLTPE GHRVAVHVYG
 201 TRQYFYMNKE EVDRHLQCRA PRDLCERMAA ALRESPGASF RGISADHFEA
 251 EVVERTDVYY YETRPALFYR VYVRSGRVLS YLCDNFCAI KKYEGGV DAT
 20 301 TRFILDNPGF VTFGWYRLKP GRNNTLAQPR APMAFGTSSD VEFNCTADNL
 351 AIEGGMSDLP AYKLMCFDIE CKAGGEDELA FPVAGHPEDL VIQISCLLYD
 25 401 LSTTALEHVL LFSLGSCDLP ESHLNELAAR GLPTPVVLEF DSEFEMLLAF
 451 MTLVKQYGPE FVTGYNIINF DWPFLAKLT DIYKVPLDGY GRMNGRGVFR
 501 VWDIGQSHFQ KRSKIKVNGM VNIDMYGIIT DKIKLSSYKL NAVA EAVLKD
 30 551 KKKDLSYRDI PAYYAAGPAQ RGVIGEYCIQ DSVLVGQLFF KFLPHLELSA
 601 VARLAGINIT RTIYDGQQIR VFTCLRLAD QKGFILPDTQ GRFRGGGGEA
 35 651 PKRPAAARED EERPEEEGED EDEREEGGGE REPEGARETA GRHVG YQGAR
 701 VLDPTSGFHV NPVVVDFDAS LYPSIIQAHN LCFSTLSLRA DAVAHLEAGK
 751 DYLEIEVGGR RLFFVKAHVR ESLLSILLRD WLAMRKQIRS RIPQSSPEEA
 40 801 VLLDKQQA AI KVCNSVYGF TGAQHGLLPC LHVAATVT TI GREMLLATRE
 851 YVHARWAAFE QLLADFPEAA DMRAPGPYSM RIYGD TDSI FVLCRGLTAA
 45 901 GLTAVGDKMA SHISRALFLS PIKLECEKTF TKLLLI AKKK YIGVTYGGKM
 951 LIKGVDLVRK NNCAFINRTS RALVDLLFYD DTVSGAAAAL AERP AEWLA

50 1001 RPLPEGLQAF GAVLVDAHRR ITDPERDIQD FVLTAELSRH PRAYTNKRLA
 1051 HLTVYYKLMA RRAQVPSIKD RPYVIVAQT REVEETVARL AALRELDAAA
 1101 PGDEFAPPAA LPSPAKRPRE TPLHADPPGG ASKPRKLLVS ELAEDPAYAI
 55 1151 AHGVALNTDY YFSHLLGAAC VTFKALFGNN AKITESLLKR FIPEVWHPPD
 1201 DVAARLRAAG FGAVGAGATA EETRRMLHRA FDTLA*

60

5 **Figure 5Q DNA and amino acid sequence list**

SEQ.ID.NO. 9 DNA sequence of HSV polymerase gene for HSV1-DJL-M1

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1 ATGTTTTCCG GTGGCGGCGG CCCGCTGTCC CCCGGAGGAA AGTCGGCGGC
10 51 CAGGGCGGCG TCCGGGTTTT TTGCGCCCGC CGGCCCTCGC GGAGCCGGCC
101 GGGGACCCCC GCCTTGTTTG AGGCAAAACT TTTACAACCC CTACCTCGCC
151 CCAGTCGGGA CGCAACAGAA GCCGACCGGG CCAACCCAGC GCCATACGTA
15 201 CTATAGCGAA TGCGATGAAT TTCGATTCAT CGCCCCGCGG GTGCTGGACG
251 AGGATGCCCC CCCGGAGAAG CGCGCCGGGG TGCACGACGG TCACCTCAAG
20 301 CGCGCCCCCA AGGTGTACTG CGGGGGGGAC GAGCGCGACG TCCTCOGCGT
351 CGGGTCGGGC GGCTTCTGGC CGCGGCGCTC GCGCCTGTGG GGCGGCGTGG
401 ACCACGCCCC GGCGGGGTTT AACCCACCG TCACCGTCTT TCACGTGTAT
25 451 GACATCCTGG AGAACGTGGA GCACGCGTAC GGCATGCGCG CGGCCAGTT
501 CCACGCGCGG TTTATGGACG CCATCACACC GACGGGGACC GTCATCACGC
30 551 TCCTGGGCCT GACTCCGGAA GGCCACCGGG TGGCCGTTCA CGTTTACGGC
601 ACGCGCAGT ACTTTTACAT GAACAAGGAG GAGGTTGACA GGCACCTACA
651 ATGCCGCGCC CCACGAGATC TCTGCGAGCG CATGGCCGCG GCCCTGCGCG
35 701 AGTCCCCGGG CGCGTCGTTT CGCGGCATCT CCGCGGACCA CTTGAGGGCG
751 GAGGTGGTGG AGCGCACCGA CGTGTACTAC TACGAGACGC GCCCCGCTCT
40 801 GTTTTACCGC GTCTACGTCC GAAGCGGGCG CGTGCTGTG TACCTGTGCG
851 ACAACTTCTG CCCGGCCATC AAGAAGTACG AGGGTGGGGT CGACGCCACC
901 ACCCGGTTCA TCCTGGACAA CCCCGGGTTC GTCACCTTCG GCTGGTACCG
45 951 TCTCAAACCG GGCCGGAACA ACACGCTAGC CCAGCCGCGG GCCCCGATGG
1001 CCTTCGGGAC ATCCAGCGAT GTCGAGTTTA ACTGTACGGC GGACAACCTG
50 1051 GCCATCGAGG GGGGCATGAG CGACCTACCG GCATACAAGC TCATGTGCTT
1101 CGATATCGAA TGCAAGGCGG GGGGGGAGGA CGAGCTGGCC TTTCGGGTGG
1151 CCGGGCACCC GGAGGACCTG GTCATCCAGA TATCCTGTCT GCTCTACGAC
55 1201 CTGTCCACCA CCGCCCTGGA GCACGTCCTC CTGTTTTCGC TCGGTTCTTG
1251 CGACCTCCCC GAATCCCACC TGAACGAGCT GGCGGCCAGG GGCTTGCCCA

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5

Figure 5R DNA and amino acid sequence list

10 1301 CGCCCGTGGT TCTGGAATTC GACAGCGAAT TCGAGATGCT GTTGGCCTTC
 1351 ATGACCCTTG TGAAACAGTA CGGCCCGAG TTCGTGACCG GGTACAACAT
 1401 AATCAACTTC GACTGGCCCT TCTTGCTGGC CAAGCTGACG GACATTTACA
 15 1451 AGGTCCCCCT GGACGGGTAC GGCOGCATGA ACGGCCGGGG CGTGTTTCGC
 1501 GTGTGGGACA TAGGCCAGAG CCACTTCCAG AAGCGCAGCA AGATAAAGGT
 20 1551 GAACGGCATG GTGAACATCG ACATGTACGG GATTATAACC GACAAGATCA
 1601 AGCTCTCGAG CTACAAGCTC AACGCCGTGG CCGAAGCCGT CCTGAAGGAC
 1651 AAGAAGAAGG ACCTGAGCTA TCGCGACATC CCCACCTACT ACGCCGCCGG
 25 1701 GCCCGCGCAA CGCGGGGTGA TCGCGAGTA CTGCATACAG GATTCCCTGC
 1751 TGGTGGGCCA GCTGTTTTTT AAGTTTTTGC CCCATCTGGA GCTCTCGGCC
 1801 GTCGCGCGCT TGGCGGGTAT TAACATCACC CGCACCATCT ACGACGGCCA
 30 1851 GCAGATCCGC GTCTTTACGT GCCTGCTGCG CCTGGCCGAC CAGAAGGGCT
 1901 TTATTCTGCC GGACACCCAG GGGCGATTTA GGGGCGCCGG GGGGGAGGCG
 35 1951 CCCAAGCGTC CGGCCGCAGC CCGGGAGGAC GAGGAGCGGC CAGAGGAGGA
 2001 GGGGGAGGAC GAGAACGAAC GCGAGGAGGG CGGGGGCGAG CGGGAGCCGG
 2051 AGGGCGCGCG GGAGACCGCC GGCCGGCAOG TGGGGTACCA GGGGGCCAGG
 40 2101 GTCCTTGACC CCACTTCCGG GTTTCACGTG AACCCCGTGG TGGTGTTCTGA
 2151 CTTTGCCAGC CTGTACCCCA GCATCATCCA GGCCCACAAC CTGTGCTTCA
 45 2201 GCACGCTCTC CCTGAGGGCC GACGCAGTGG CGCACCTGGA GGCGGGCAAG
 2251 GACTACCTGG AGATCGAGGT GGGGGGGCGA CGGCTGTTCT TCGTCAAGGC

 50 2301 TCACGTGCGA GAGAGCCTCC TCAGCATCCT CCTGCGGGAC TGGCTCGCCA
 2351 TGCGAAAGCA GATCCGCTCG CGGATTCCCC AGAGCAGCCC CGAGGAGGCC
 2401 GTGCTCCTGG ACAAGCAGCA GGCCGCCATC AAGGTCGTGT GTAACTCGGT
 55 2451 TTACGGGTTC ACGGGAGCGC AGCACGGACT CCTGCCGTGC CTGCACGTTG
 2501 CCGCGACGGT GACGACCATC GGCCGCGAGA TGCTGCTCGC GACCCGCGAG
 2551 TACGTCCACG CGCGCTGGGC GGCCTTCGAA CAGCTCCTGG CCGATTTCCT

60

5 **Figure 5S DNA and amino acid sequence list**

2601 GGAGGCGGCC GACATGCGCG CCCCCGGGCC CTATTCCATG CGCATCATCT
 10 2651 ACGGGGACAC GGACTCCATA TTTGTGCTGT GCCGCGGCCT CACGGCCGCC
 2701 GGGCTGACGG CCGTGGGCGA CAAGATGGCG AGCCACATCT CGCGCGCGCT
 2751 GTTCTGCCC CCCATCAAAC TCGAGTGCGA AAAGACGTTC ACCAAGCTGC
 15 2801 TGCTGATCGC CAAGAAAAAG TACATCGGCG TCATCTACGG GGGTAAGATG
 2851 CTCATCAAGG GCGTGGATCT GGTGCGCAA AACAAGTGG CGTTTATCAA
 20 2901 CCGCACCTCC AGGGCCCTGG TCGACCTGCT GTTTTACGAC GATACCGTAT
 2951 CCGGAGCGGC CGCCGCGTTA GCCGAGCGCC CCGCAGAGGA GTGGCTGGCG
 3001 CGACCCCTGC CCGAGGGACT GCAGGCGTTC GGGGCGGTCC TCGTAGACGC
 25 3051 CCATCGGCGC ATCACCGACC CGGAGAGGGA CATCCAGGAC TTTGTTCTCA
 3101 CCGCCGA ACT GAGCAGACAC CCGCGCGCGT ACACCAACAA GCGCCTGGCC
 30 3151 CACCTGACGG TGTATTACAA GTCATGGCC CGCCGCGCGC AGGTCCCGTC
 3201 CATCAAGGAC CGGATCCCGT ACGTGATCGT GGCCCAGACC CGCGAGGTAG
 3251 AGGAGACGGT CGCGCGGCTG GCGCCCTCC GCGAGCTAGA CGCCGCCGCC
 35 3301 CCAGGGGACG AGCCCGCCCC CCGCGGGCC CTGCCCTCCC CGGCCAAGCG
 3351 CCCCCGGGAG ACGCCGTCGC CTGCCGACCC CCGGGAGGC GCGTCCAAGC
 40 3401 CCCGCAAGCT GCTGGTGTCC GAGCTGGCCG AGGATCCCGC ATACGCCATT
 3451 GCCACGGCG TCGCCCTGAA CACGGA CTAT TACTTCTCCC ACCTGTTGGG
 3501 GCGGCGTGC GTGACATTCA AGGCCCTGTT TGGGAATAAC GCCAAGATCA
 45 3551 CCGAGAGTCT GTTAAAAAGG TTTATTCCCG AAGTGTGGCA CCCCCGGAC
 3601 GACGTGGCCG CGCGGCTCCG GACCGCAGGG TTCGGGGCGG TGGGTGCCGG
 50 3651 CGCTACGGCG GAGGAACTC GTCGAATGTT GCATAGAGCC TTTGATACTC
 3701 TAGCATGA

5 **Figure 5T DNA and amino acid sequence list****SEQ.ID.NO. 10** Amino acid sequence of DNA polymerase for HSV1-DJL-M1

1 MFSGGGGPLS PGGKSAARAA SGFFAPAGPR GAGRGPPPCL RQNFYNPYLA
 10 51 PVGTQKQPTG PTQRHTYYSE CDEFRFIAPR VLDEDAPEEK RAGVHDGHLK
 101 RAPKVYCGGD ERDVLRVGSG GFWPRRSRLW GGVDHAPAGF NPTVTVFHVY
 15 151 DILENVEHAY GMRAAQFHAR FMDAITPTGT VITLLGLTPE GHRVAVHVY
 201 TRQYFYMNKE EVDRHLQCRA PRDLCERMAA ALRESPGASF RGISADHFEA
 251 EVVERTDVYY YETRPALFYR VYVRSGRVLS YLCDNFCPAI KKYEGGVDAT
 20 301 TRFILDNPGF VTFGWYRLKP GRNNTLAQPR APMAFGTSSD VEFNCTADNL
 351 AIEGGMSDLP AYKLMCFDIE CKAGGEDELA FPVAGHPEDL VIQISCLLYD
 401 LSTTALEHVL LFSLGSCDLP ESHLNELAAR GLPTPVVLEF DSEFEMLLAF
 25 451 MTLVKQYGPE FVTGYNIINF DWPFLAKLT DIYKVPLDGY GRMNGRGVFR
 501 VWDIGQSHFQ KRSKIKVNGM VNIDMYGIIT DKIKLSSYKL NAVAEAVLKD
 30 551 KKKDLSYRDI PTYYAAGPAQ RGVIGEYCIQ DSSLVGQLFF KFLPHLELSA
 601 VARLAGINIT RTIYDGQQIR VFTCLRLAD QKGFILPDTQ GRFRGAGGEA
 651 PKRPAAARED EERPEEEGED ENEREEGGGE REPEGARETA GRHVGYQGAR
 35 701 VLDPTSGFHV NPVVVFDFAS LYPSIIQAHN LCFSTLSLRA DAVAHLEAGK
 751 DYLEIEVGGR RLFFVKAHVR ESLLSILLRD WLAMRKQIRS RIPQSSPEEA
 40 801 VLLDKQQAII KVV CNSVYGF TGAQHGLLPC LHVAATVTTI GREMLLATRE
 851 YVHARWAAFE QLLADFPEAA DMRAPGPYSM RIYGD TDSI FVLCRGLTAA
 901 GLTAVGDKMA SHISRALFLP PIKLECEKTF TKLLLIAKKK YIGVITYGGKM
 45 951 LIKGVDLVRK NNCAFINRTS RALVDLLFYD DTVSGAAAAL AERPAAEWLA
 1001 RPLPEGLQAF GAVLVDAHRR ITDPERDIQD FVLTAELSRH PRAYTNKRLA

 50 1051 HLTVYYKLMA RRAQVPSIKD RPYVIVAQT REVEETVARL AALRELDAAA
 1101 PGDEPAPPAA LPSPAKRPRE TPSPADPPGG ASKPRKLLVS ELAEDPAYAI
 1151 AHGVALNTDY YFSHLLGAAC VTFKALFGNN AKITESLLKR FIPEVWHPPD
 55 1201 DVAARLRTAG FGAVGAGATA EETRRMLHRA FDTLA*

5 **Figure 5U DNA and amino acid sequence list**

SEQ.ID.NO. 11 DNA sequence of DNA polymerase gene for HMCV-AD169-M1

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10   1 ATGTTTTTCA ACCCGTATCT GAGCGGCGGC GTGACCGGCG GTGCGGTGCG
    51 GGGTGGCCGG CGTCAGCGTT CGCAGCCCGG CTCCGCGCAG GGCTCGGGCA
    101 AGCGGCCGCC ACAGAAACAG TTTTGCAGA TCGTGCCGCG AGGTGTCATG
15   151 TTCGACGGTC AGACGGGGTT GATCAAGCAT AAGACGGGAC GGCTGCCTCT
    201 CATGTTCTAT CGAGAGATTA AACATTGTT GAGTCATGAC ATGGTTTGGC
    251 CGTGTCTTG GCGCGAGACC CTGGTGGGTC GCGTGGTGGG ACCTATTCGT
    301 TTTCACACCT ACGATCAGAC GGACGCCGTG CTCTTCTTCG ACTCGCCCGA
    351 AAACGTGTCG CCGCGCTATC GTCAGCATCT GGTGCCTTCG GGGAACGTGT
25   401 TCGGTTTCTT CGGGGCCACA GAACACGGCT ACAGTATCTG CGTCAACGTT
    451 TTCGGGCAGC GCAGCTACTT TTA CTGTGAG TACAGCGACA CCGATAGGCT
    501 GCGTGAGGTC ATTGCCAGCG TGGGCGAACT AGTGCCCGAA CCGCGGACGC
    551 CATA CGCCGT GTCTGTCACG CCGGCCACCA AGACCTCCAT CTATGGGTAC
    601 GGGACGCGAC CCGTGCCCGA TTG CAGTGT GTGTCTATCA GCAACTGGAC
35   651 CATGGCCAGA AAAATCGGCG AGTATCTGCT GGAGCAGGGT TTTCCCGTGT
    701 ACGAGGTCCG TGTGGATCCG CTGACGCGTT TGGTCATCGA TCGGCGGATC
    751 ACCACGTTCG GCTGGTGCTC CGTGAATCGT TACGACTGGC GGCAGCAGGG
    801 TCGCGCGTCG ACTTGTGATA TCGAGGTAGA CTGCGATGTC TCTGACCTGG
    851 TGGCTGTGCC CGACGACAGC TCGTGGCCGC GCTATCGATG CCTGTCCTTC
45   901 GATATCGAGT GCATGAGCGG CGAGGGTGGT TTTCCCTGCG CCGAGAAGTC
    951 CGATGACATT GTCATT CAGA TCTCGTGCGT GTGCTACGAG ACGGGGGGAA
50  1001 ACACCGCCGT GGATCAGGGG ATCCCAAACG GGAACGATGG TCGGGGCTGC
    1051 ACTTCGGAGG GTGTGATCTT TGGGCACTCG GGTCTTCATC TCTTTACGAT
    1101 CGGCACCTGC GGGCAGGTGG GCCCAGACGT GGACGTCTAC GAGTTCCTT
55  1151 CCGAATACGA GCTGCTGCTG GGCTTTATGC TTTTCTTCA ACGGTACGCG
    1201 CCGGCCTTTG TGACCGGTTA CAACATCAAC TCTTTTGACT TGAAGTACAT
60

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5 **Figure 5V DNA and amino acid sequence list**

1251 CCTCACGCGT CTCGAGTACC TGTATAAGGT GGACTCGCAG CGCTTCTGCA
 10 1301 AGTTGCCTAC GGCGCAGGGC GGCCGTTTCT TTTTACACAG CCCCGCCGTG
 1351 GGTTTTAAGC GGCAGTACGC CGCCGCTTTT CCCTCGGCTT CTCACAACAA
 1401 TCCGGCCAGC ACGGCCGCCA CCAAGGTGTA TATTGCGGGT TCGGTGGTTA
 15 1451 TCGACATGTA CCCTGTATGC ATGGCCAAGA CTAAC TCGCC CAACTATAAG
 1501 CTCAACACTA TGGCCGAGCT TTACCTGCGG CAACGCAAGG ATGACCTGTC
 20 1551 TTACAAGGAC ATCCCGCGTT GTTTCGTGGC TAATGCCGAG GGCCGCGCCC
 1601 AGGTAGGCCG TTAGTGTCTG CAGGACGCCG TATTGGTGCG CGATCTGTTC
 1651 AACACCATTA ATTTTCACTA CGAGGCCGGG GCCATCGCGC GGCTGGCTAA
 25 1701 AATTCCGTTG CGGCGTGTCA TCTTTGACGG ACAGCAGATC CGTATCTACA
 1751 CCTCGCTGCT GGACGAGTGC GCCTGCCGCG ATTTTATCCT GCCCAACCAC
 30 1801 TACAGCAAAG GTACGACGGT GCCCGAAACG AATAGCGTTG CTGTGTCACC
 1851 TAACGCTGCT ATCATCTCTA CCGCCGCTGT GCCCGGCGAC GCGGGTTCTG
 1901 TGGCGGCTAT GTTTCAGATG TCGCCGCCCT TGCAATCTGC GCCGTCCAGT
 35 1951 CAGGACGGCG TTTCACCCGG CTCCGGCAGT AACAGTAGTA GCAGCGTCGG
 2001 CGTTTTTCAGC GTCGGCTCCG GCAGTAGTGG CGGCGTCGGC GTTCCAACG
 40 2051 ACAATCACGG CGCCGGCGGT ACTGCGGCGG TTTCGTACCA GGGCGCCACG
 2101 GTGTTTGAGC CCGAGGTGGG TTAGTACAAC GACCCCGTGG CCGTGTTTGA
 2151 CTTTGCCAGC CTCTACCCTT CCATCATCAT GGCCCACAAC CTCTGCTACT
 45 2201 CCACCCTGCT GGTGCCGGGT GGCGAGTACC CTGTGGACCC CGCCGACGTA
 2251 TACAGCGTCA CGCTAGAGAA CGGCGTGACC CACCGCTTTG TCGTGCTTC
 50 2301 GGTGCGCGTC TCGGTGCTCT CGGAAGTGT CAACAAGTGG GTTTCGCAGC
 2351 GGCGTGCCGT GCGCGAATGC ATGCGCGAGT GTCAAGACCC TGTGCGCCGT
 2401 ATGCTGCTCG ACAAGGAACA GATGGCGCTC AAAGTAACGT GCAACGCTTT
 55 2451 CTACGGTTTT ACCGGCGCGC TGAACGGTAT GATGCCGTGT CTGCCCATCG
 2501 CCGCCAGCAT CACGCGCATC GGTCGCGACA TGCTAGAGCG CACGGCGCGG

5

Figure 5W DNA and amin acid sequence list

10 2551 TTCATCAAAG ACAACTTTTC AGAGCCGTGT TTTTGCACA ATTTTTTTAA
 2601 TCAGGAAGAC TATGTAGTGG GAACGCGGGA GGGGGATTCG GAGGAGAGCA
 2651 GCGCGTTACC GGAGGGGCTC GAAACATCGT CAGGGGGCTC GAACGAACGG
 15 2701 CGGGTGGAGG CGCGGGTCAT CTACGGGGAC ACGGACAGCG TGTTTGTCCG
 2751 CTTTCGTGGC CTGACGCCGC AGGCTCTGGT GGCGCGTGGG CCCAGCCTGG
 20 2801 CGCACTACGT GACGGCCTGT CTTTTTGTGG AGCCCGTCAA GCTGGAGTTT
 2851 GAAAAGGTCT TCGTCTCTCT TATGATGATC TGCAAGAAAC GTTACATCGG
 2901 CAAAGTGGAG GGCGCCTCGG GTCTGAGCAT GAAGGGCGTG GATCTGGTGC
 25 2951 GCAAGACGGC CTGCGAGTTC GTCAAGGGCG TCACGCGTGA CGTCCTCTCG
 3001 CTGCTCTTTG AGGATCGCGA GGTCTCGGAA GCAGCCGTGC GCCTGTCCGG
 30 3051 CCTCTCACTC GATGAAGTCA AGAAGTACGG CGTGCCACGC GGTTCCTGGC
 3101 GTATCTTACG CCGCTTGGTG CAGGCCCGCG ACGATCTGTA CCTGCACCGT
 3151 GTGCGTGTCG AGGACCTGGT GCTTTCGTCTG GTGCTCTCTA AGGACATCTC
 35 3201 GCTGTACCGT CAATCTAACC TGCCGCACAT TGCCGTCATT AAGCGATTGG
 3251 CGGCCCCGTT TGAGGAGCTA CCCTCGGTCT GGGATCGGGT CTTTACGTT
 40 3301 CTGACGGCGC CCGGTGTCCG GACGGCGCGC CAGGGTTCCT CCGACAACGG
 3351 TGATTCTGTA ACCGCCGGCG TGGTTTCCCG GTCGGACGCG ATTGATGGCA
 3401 CGGACGACGA CGCTGACGGC GGCGGGGTAG AGGAGAGCAA CAGGAGAGGA
 45 3451 GGAGAGCCGG CAAAGAAGAG GGCGCGGAAA CCACCGTCGG CCGTGTGCAA
 3501 CTACGAGGTA GCCGAAGATC CGAGCTACGT GCGCGAGCAC GGCGTGCCCA
 50 3551 TTCACGCCGA CAAGTACTTT GAGCAGGTTC TCAAGGCTGT AACTAACGTG
 3601 CTGTCGCCCC TCTTCCCGG CGGCGAAACC GCGCGCAAGG ACAAGTTTTT
 3651 GCACATGGTG CTGCCGCGGC GCTTGCACTT GGAGCCGGCT TTTCTGCCGT
 55 3701 ACAGTGTCAA GGCGCACGAA TGCTGTTGA

5 **Figure 5X DNA and amino acid sequence list****SEQ.ID.NO.12 Amino acid sequence of DNA polymerase f r HCMV-AD169-M1**

10 1 MFFNPYLSGG VTGGAVAGGR RQRSQPGSAQ GSGKRPPQKQ FLQIVPRGVM
 51 FDGQTGLIKH KTGRLPLMFY REIKHLLSHD MVWPCPWRET LVGRVVGPIR
 101 FHTYDQTDV LFFDSPENV S PRYRQHLVPS GNVLRFFGAT EHGYSICNVV
 15 151 FGQRSYFYCE YSDTDLREV IASVGELVPE PRTPYAVSVT PATKTSIYGY
 201 GTRPVFDLQC VSIWNWTMAR KIGEYLLQGG FPVYEVRVDP LTRLVIDRRI
 20 251 TTFGWCSVNR YDWRQQGRAS TCDIEVDCDV SDLVAVPDDS SWPRYRCLSF
 301 DIECMSGEGG FPCAESDDI VIQISCVCYE TGGNTAVDQG IPNGNDGRGC
 351 TSEGVIFGHS GLHLFTIGTC GQVGPDVDVY EFPSEYELL GFMLFFQRYA
 25 401 PAFVTGYNN SFDLKYILTR LEYLYKVDSQ RFCKLPTAQG GRFFLHSPAV
 451 GFKRQYAAAF PSASHNNPAS TAATKVYIAG SVVIDMYPVC MAKTNSPNYK
 30 501 LNTMAELYLR QRKDDLSYKD IPRCFVANAE GRAQVGRYCL QDAVLVRDLF
 551 NTINFHYEAG AIARLAKIPL RRVIFDGQOI RIYTSLLDEC ACRDFILPNH
 601 YSKGTTVPET NSVAVSPNAA IISTAAPGD AGSVAAMFQM SPPLQSAPSS
 35 651 QDGVSPGSGS NSSSSVGVFS VSGSGSSGGVG VSNDNHGAGG TAAVSYQGAT
 701 VFEPEVGYYN DPVAVFDFAS LYPSTMAHN LCYSTLLVPG GEYPVDPADV
 40 751 YSVTLENGVT HRFVRASVRV SVLSELLNKW VSQRRVREC MRECQDPVRR
 801 MLLDKEQMAL KVTCTAFYGF TGALNGMMPC LPIAASITRI GRDMLERTAR
 851 FIKDNFSEPC FLHNFFNQED YVVGTTREGDS EESSALPEGL ETSSGGSNER
 45 901 RVEARVIYGD TDSVFVRFRG LTPQALVARG PSLAHYVTAC LFVEPVKLEF
 951 EKVFVSLMMI CKKRYIGKVE GASGLSMKGV DLVRKTACEF VKGVTRDVLV
 50 1001 LFFEDREVSE AAVRLSRLSL DEVKKYGVPR GFWRILRRLV QARDDLYLHR
 1051 VRVEDLVLS VLSKDISLYR QSNLPHIAVI KRLAARSEEL PSVGDRVFYV
 1101 LTAPGVRTAP QGSSDNGDSV TAGVVSRS DA IDGTDDDADG GGVEESNRRG
 55 1151 GEPAKKRARK PPSAVCNYEV AEDPSYVREH GVPIHADKYF EQVLKAVTNV
 1201 LSPVFPGET ARKDKFLHMV LPRRLHLEPA FLPYSVKAHE CC*